



STIC Search Report

Biotech-Chem Library

STIC Database Tracking Number: 176925

TO: James Schultz
Location: 2d18 / 2c18
Art Unit: 1635 *2/c*
Monday, January 30, 2006

Case Serial Number: 09/227881

From: Noble Jarrell
Location: Biotech-Chem Library
Rem 1B71
Phone: 272-2556

Noble.jarrell@uspto.gov

Search Notes

STIC-Biotech/ChemLib

176925

From: Schultz, James
Sent: Wednesday, January 18, 2006 3:12 PM
To: STIC-Biotech/ChemLib
Subject: Seq Search 09/227,881

Hello,

Could you please run a score over length sequence search on SEQ ID NO:34 (5271 nt long), which returns hits between 50 and 500 nt long? I need only those hits that are 100% complementary. Please run the search in the interference databases as well.

Also, could you run a standard nucleotide sequence search on the full length of SEQ ID NO: 34, ONLY in the interference databases?

Thanks,
Doug Schultz

James Douglas Schultz, PhD
Primary Examiner
AU 1635 (Biotechnology)
United States Patent and Trademark Office
(Office) REM 2D18
(Mail) REM 2C18
(571) 272-0763

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Searcher: ndle
Searcher Phone: _____
Date Searcher Picked up: 1/18/06
Date completed: _____
Searcher Prep Time: 45
Online Time: 10

Type of Search
NA# 11 AA# 1
S/L: ✓ Oligomer: ✓
Encode/Transl: _____
Structure #: _____ Text: _____
Inventor: _____ Litigation: _____

Vendors and cost where applicable
STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: CompuGen
WWW/Internet: _____
Other (Specify): gcg

1	5271	100.0	5271	3	US-09-306-828-34	Sequence 34, App
2	5271	100.0	6169	3	US-08-938-669A-3	Sequence 3, Appl
3	5271	100.0	6169	3	US-09-306-828-3	Sequence 3, Appl
4	5246.4	99.5	5300	3	US-08-938-669A-1	Sequence 1, Appl
5	5246.4	99.5	5300	3	US-09-306-828-1	Sequence 1, Appl
6	5224.4	99.1	5304	3	US-08-938-669A-2	Sequence 2, Appl
7	5224.4	99.1	5304	3	US-09-306-828-2	Sequence 2, Appl
8	1804.4	34.2	2800	3	US-09-056-828A-1	Sequence 1, Appl
9	1804.4	34.2	2800	3	US-10-017-870-10	Sequence 10, Appl
10	1804.4	34.2	2800	3	US-09-552-464A-1	Sequence 1, Appl
11	227	4.3	227	3	US-09-306-828-38	Sequence 38, App
12	227	4.3	283	3	US-09-306-828-37	Sequence 37, App
13	189.8	3.6	29326	3	US-09-949-016-15556	Sequence 15356, App
14	189.8	3.5	205044	3	US-09-949-016-15851	Sequence 15851, App
15	184.6	3.5	205044	3	US-09-949-016-15852	Sequence 15852, App
16	184.6	3.5	205044	3	US-09-949-016-15853	Sequence 15853, App
17	184.6	3.5	223471	3	US-09-949-016-12187	Sequence 12187, App
18	184.6	3.5	223471	3	US-09-949-016-12724	Sequence 12724, App
19	184.6	3.5	223471	3	US-09-949-016-12725	Sequence 12725, App
20	184.4	3.5	1744	3	US-10-104-047-1845	Sequence 1845, App
21	183.2	3.5	88906	3	US-09-949-016-17668	Sequence 17468, App
22	183	3.5	113966	3	US-09-949-016-12277	Sequence 12277, App
23	183	3.5	113967	3	US-09-949-016-17051	Sequence 17051, App
24	182.8	3.5	601	3	US-09-949-016-10621	Sequence 10621, App

C	25	182.8	3.5	601	3	US-09-949-016-102733	Sequence 102733
C	26	182.8	3.5	76164	3	US-09-949-016-12288	Sequence 12288
C	27	182.8	3.5	76165	3	US-09-949-016-14005	Sequence 14005
C	28	182.6	3.5	601	3	US-09-949-016-185608	Sequence 185608
C	29	181.6	3.4	678633	3	US-09-949-016-14577	Sequence 14577
C	30	181.6	3.4	678633	3	US-09-949-016-14578	Sequence 14578
C	31	181.2	3.4	360470	3	US-09-949-016-11173	Sequence 13173
C	32	181	3.4	601	3	US-09-949-016-185607	Sequence 185607
C	33	181	3.4	12361	3	US-09-949-016-16870	Sequence 16870
C	34	181	3.4	44120	3	US-09-949-016-11151	Sequence 14151
C	35	181	3.4	44120	3	US-09-949-016-11152	Sequence 14152
C	36	181	3.4	44120	3	US-09-949-016-11153	Sequence 14153
C	37	181	3.4	44120	3	US-09-949-016-11154	Sequence 14154
C	38	181	3.4	44120	3	US-09-949-016-11155	Sequence 14155
C	39	181	3.4	44120	3	US-09-949-016-14156	Sequence 14156
C	40	181	3.4	265030	3	US-09-949-016-15779	Sequence 15779
C	41	180.8	3.4	601	3	US-09-949-016-19567	Sequence 19567
C	42	180.8	3.4	601	3	US-09-949-016-117194	Sequence 117194
C	43	180.6	3.4	66933	3	US-09-544-398B-11	Sequence 11, Appl
C	44	180.6	3.4	66933	3	US-09-544-398B-11	Sequence 11, Appl
C	45	180.6	3.4	72049	3	US-09-544-398B-9	Sequence 9, Appl

ALIGNMENTS

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RESULT 1
US-09-306-828-34
; Sequence 34, Application US/09306828
; Patent No. 6475724
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis, Prognosis
; CURRENT APPLICATION NUMBER: US/09/306,828
; EARLIER FILING DATE: 1999-05-07
; EARLIER APPLICATION NUMBER: US 09/227,881
; EARLIER FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 34
; LENGTH: 5271
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-306-828-34,

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Query Match	Score	DB 3	Length
100.0%	5271		5271

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Best Local Similarity    100.0%;   Pred. NO. 0;
Matches 5271; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Oy	121	GGAAGAGAGATATCCAGTTTAGCCAAATGTTCAGGCTGTGTCTGCTTTATTTTAAGTA	180
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Oy	181	CAGATTTTCTCTCTGACAGAGCTTTTCTTACAGGAAACATCACATTCACAATATGTGAATC	240
Db	181	CAGATTTTCTCTCTGACAGAGCTTTTCTTACAGGAAACATCACATTCACAATATGTGAATC	240
Oy	241	CATCAACACAGAGCTTAAGAAACAGGAATGGAATGGGCACTGGCCACAGAAAAATGCGAG	3000
Db	241	CATCAACACAGAGCTTAAGAAACAGGAATGGAATGGGCACTGGCCACAGAAAAATGCGAG	3000
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 Db 421 TAATTAAGTATTTTCTTGGGAGAGACCTCCATGTGAGCTTGATGGAAAAATGGAA 480
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QY 3541 GTTCTAG 3600
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Db 3661 GTAGTAACTGAGAGCTGATAGATTTACTAGTTTCTCTTATTAAGAACTCTTTTCTGTGT 3720
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Db 4621 CATGACACACAGAGATTAAG 4680
QY 4681 TGCAGAGCTGAATTAAG 4740

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Db	841	GGGACCCCTGAGGCATTTCCCTTTAGGAAGGCCAGTTTTCTTAAGGAATCTTTAAGAAATCTC	900
Qy	901	TTGAAAGATCATGAATTTTAAACCAATTTTAAAGTATAAAACAAATATGCGATGCAATATCAG	960
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Qy	961	TTTATAGCATGGTCCCAATTTTATAAAGTCAGGCATACAAGGATAACGTCGCCAGCTCC	1020
Db	961	TTTATAGCATGGTCCCAATTTTATAAAGTCAGGCATACAAGGATAACGTCGCCAGCTCC	1020
Qy	1021	GGATAGGTCAGAAATCATTTAGAAATCACTGTGTCCCACTCTAAGCTTTTTCAGAAATGATC	1080
Db	1021	GGATAGGTCAGAAATCATTTAGAAATCACTGTGTCCCACTCTAAGCTTTTTCAGAAATGATC	1080
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Qy	1141	GTGCTCAACCAATTTTAAAGTCATCTCAGTAGGTCCTCAATTAACAATGCCACCTCC	1200
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Qy	1261	TACAGCAGAAAGTCCCGTGAAGGTCTGTGTCTTACCACTACCTGATGCTCTAC	1320
Db	1261	TACAGCAGAAAGTCCCGTGAAGGTCTGTGTCTTACCACTACCTGATGCTCTAC	1320
Qy	1321	ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGCTCAGCCTCC	1380
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Db	1381	CGCGTAGCTGGGACTACAGGCGCACGCGCGCTAAATTTTGTATTTGTAGTAGAGATGGG	1440
Qy	1441	GTTCACATATTAAGCCGCTGTCTTGAACTCTGACCTCAGGTATCACCCACTC	1500
Db	1441	GTTCACATATTAAGCCGCTGTCTTGAACTCTGACCTCAGGTATCACCCACTC	1500
Qy	1501	AGCTCTCAAGTCTGGGATTAAGGATGAGTCAAGGCGCGCGCTGAGTCAAGGTCAAGT	1560
Db	1501	AGCTCTCAAGTCTGGGATTAAGGATGAGTCAAGGCGCGCGCTGAGTCAAGGTCAAGT	1560
Qy	1561	TTAATAAGGAATACTTGAATGGTTTAAACCAACAGGGAACAGACAAAGCTGTGA	1620
Db	1561	TTAATAAGGAATACTTGAATGGTTTAAACCAACAGGGAACAGACAAAGCTGTGA	1620
Qy	1621	TAATTCAGGATTTCTTGGGATGGGAAATGGTGCATGAGCTGCTGCTAGTCCAGAC	1680
Db	1621	TAATTCAGGATTTCTTGGGATGGGAAATGGTGCATGAGCTGCTGCTAGTCCAGAC	1680
Qy	1681	CAGTGGCTCATCATCTTCTCCCTCATCTCATTTTTCAGGCTAAGTTACCAATTTTAT	1740
Db	1681	CAGTGGCTCATCATCTTCTCCCTCATCTCATTTTTCAGGCTAAGTTACCAATTTTAT	1740
Qy	1741	CACCATGCTTTTGTGGTAAGCTCCACATCGTTACTGAAATTAAGAGTATACATAAAGCTAG	1800
Db	1741	CACCATGCTTTTGTGGTAAGCTCCACATCGTTACTGAAATTAAGAGTATACATAAAGCTAG	1800
Qy	1801	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGAGGGGCAATACCCAGAGACTCCT	1860
Db	1801	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGAGGGGCAATACCCAGAGACTCCT	1860
Qy	1861	TGAAGCCCCGGGAGAGTTTCTCTCAGCTGGGAGCCCTGCAAGACCCGGGTCC	1920
Db	1861	TGAAGCCCCGGGAGAGTTTCTCTCAGCTGGGAGCCCTGCAAGACCCGGGTCC	1920
Qy	1921	TGGGTGTCTTACAGCAACCTGCGCCGCTGCCACTGGTTGTGTTTATCACTCTCTAGG	1980
Db	1921	TGGGTGTCTTACAGCAACCTGCGCCGCTGCCACTGGTTGTGTTTATCACTCTCTAGG	1980
Qy	1981	GACCTGTGCTTTCTTCTATTTCTGTGTGACTGTGTTCAATTCACAGGCATTTATTGACAAAT	2040
Db	1981	GACCTGTGCTTTCTTCTATTTCTGTGTGACTGTGTTCAATTCACAGGCATTTATTGACAAAT	2040
Qy	2041	TATTGAGTACTTATCTGCGCAGACACAGAGACAAATATGTTAGCAAGCAGTCACTGC	2100
Db	2041	TATTGAGTACTTATCTGCGCAGACACAGAGACAAATATGTTAGCAAGCAGTCACTGC	2100
Qy	2101	CTTACCTCTGTTGAGGTGACAGTTTCTCATGGAAAGCTGCAAGAAATAATTAAGCCA	2160
Db	2101	CTTACCTCTGTTGAGGTGACAGTTTCTCATGGAAAGCTGCAAGAAATAATTAAGCCA	2160
Qy	2161	GCCAACTTAAACCCAGTCTGAAAGAAAGAAATAAACCAATCTTGAAGAAATTTGTGGC	2220
Db	2161	GCCAACTTAAACCCAGTCTGAAAGAAAGAAATAAACCAATCTTGAAGAAATTTGTGGC	2220
Qy	2221	AGCATCCCTTAAACAGGCCACCTCCCTAGCGCCCTGCTGCTTCCATCGTGCCTCGGAGG	2280
Db	2221	AGCATCCCTTAAACAGGCCACCTCCCTAGCGCCCTGCTGCTTCCATCGTGCCTCGGAGG	2280
Qy	2281	CCCCAAGCCGAGTCTTCCAAAGCCTCTCTCTCATCAGTCAAGCGCTGCGCTGGCCT	2340
Db	2281	CCCCAAGCCGAGTCTTCCAAAGCCTCTCTCTCATCAGTCAAGCGCTGCGCTGGCCT	2340
Qy	2341	GCCTCGCTTCCGCTGAATCTGCTGTGTGATCTGAGCTGAGACTCTTGGCTCCAGGCT	2400
Db	2341	GCCTCGCTTCCGCTGAATCTGCTGTGTGATCTGAGCTGAGACTCTTGGCTCCAGGCT	2400
Qy	2401	CCAGAAAGAAATGGAGAGGAACTAGTCTTAAACGAGAAATCTGAGGGGACAGTGTTC	2460
Db	2401	CCAGAAAGAAATGGAGAGGAACTAGTCTTAAACGAGAAATCTGAGGGGACAGTGTTC	2460
Qy	2461	CTCAGAGGAAAGGGGCTCCACGTCAGAGGAAATTCAGAGAGTGGGGAGCTGAGGGAG	2520
Db	2461	CTCAGAGGAAAGGGGCTCCACGTCAGAGGAAATTCAGAGAGTGGGGAGCTGAGGGAG	2520
Qy	2521	TGGGAGCTGGGGCTGAGCGGTGCTGAAAGGAGGAGGAGGTAAGGAGGAGGCTGAA	2580
Db	2521	TGGGAGCTGGGGCTGAGCGGTGCTGAAAGGAGGAGGAGGTAAGGAGGAGGCTGAA	2580
Qy	2581	GCTGCCAGATGTTTCAAGTGTGTTTACGGGGCTGGGAGTTTTCCTGTTCTCTGTCAGC	2640
Db	2581	GCTGCCAGATGTTTCAAGTGTGTTTACGGGGCTGGGAGTTTTCCTGTTCTCTGTCAGC	2640
Qy	2641	CTTTTATCTTTCTCTGCTTGGAGAGAAAGTCTATTTCAAGAGGAGTGCAGTTTC	2700
Db	2641	CTTTTATCTTTCTCTGCTTGGAGAGAAAGTCTATTTCAAGAGGAGTGCAGTTTC	2700
Qy	2701	ATAAGTCAGCTGTTAAATTTCCAGGTGTGATGGGTTTCTTCCACGAGGCTTTAT	2760
Db	2701	ATAAGTCAGCTGTTAAATTTCCAGGTGTGATGGGTTTCTTCCACGAGGCTTTAT	2760
Qy	2761	TTAATGGGAATATAGGAAGCGAGCTCAATTTCTTAGGCGGTTAAATTCAGGAAGAGTGAC	2820
Db	2761	TTAATGGGAATATAGGAAGCGAGCTCAATTTCTTAGGCGGTTAAATTCAGGAAGAGTGAC	2820
Qy	2821	TGAGTCTTTTCTTTTCTTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTT	2880
Db	2821	TGAGTCTTTTCTTTTCTTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTT	2880
Qy	2881	TGCAAGACGCTCGAAGAACCTTGGAAATCAGGAGACTCGGTTTCTTTCTTTCTTTCTTTCTTT	2940
Db	2881	TGCAAGACGCTCGAAGAACCTTGGAAATCAGGAGACTCGGTTTCTTTCTTTCTTTCTTTCTTT	2940
Qy	2941	GGTGGCTGTGGCAGCGTGGCAAGTGTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT	3000
Db	2941	GGTGGCTGTGGCAGCGTGGCAAGTGTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT	3000
Qy	3001	ATAAGACCCCTTGCAGCTCTCGTGTCTGTGAAACATCTCCCTGTGATTTCTCTGTGAGGG	3060
Db	3001	ATAAGACCCCTTGCAGCTCTCGTGTCTGTGAAACATCTCCCTGTGATTTCTCTGTGAGGG	3060

QY	1741	CACCATGCTTTTGTGTAAAGCTCCACATCGTTACTGAAATAGAGTATATACATAAAGTAG	1800	QY	2821	TGGAGTCTTTTCTTTTCATGTCCTTCTGGGCAACTACTCAGCCCTGTGGTGGACTTGGCTTA	2880
DB	1741	CACCATGCTTTTGTGTAAAGCTCCACATCGTTACTGAAATAGAGTATATACATAAAGTAG	1800	DB	2821	TGGAGTCTTTTCTTTTCATGTCCTTCTGGGCAACTACTCAGCCCTGTGGTGGACTTGGCTTA	2880
QY	1801	TTCCATTTGGGGCCATCTGTGTGTGTGTATAGGGAGGGGATACCCCCAGAGACTCTCT	1860	QY	2881	TCGAAGACGGTTCGAAAACTTGGAAATCAGGAGATCGGGTTTCTTCTGGTTCGCAAT	2940
DB	1801	TTCCATTTGGGGCCATCTGTGTGTGTGTATAGGGAGGGGATACCCCCAGAGACTCTCT	1860	DB	2881	TCGAAGACGGTTCGAAAACTTGGAAATCAGGAGATCGGGTTTCTTCTGGTTCGCAAT	2940
QY	1861	TGAAGCCCGGGCAGAGGTTTCTCTCAGCTGGGGAGCCCTGCAAGCAACCCGGGGTCC	1920	QY	2941	GGTTGGCTGTGGACCGGTGGCAAGTGTCTCTCCCTTCCCTGGGCCATAGTCTCTCTGCT	3000
DB	1861	TGAAGCCCGGGCAGAGGTTTCTCTCAGCTGGGGAGCCCTGCAAGCAACCCGGGGTCC	1920	DB	2941	GGTTGGCTGTGGACCGGTGGCAAGTGTCTCTCCCTTCCCTGGGCCATAGTCTCTCTGCT	3000
QY	1921	TGGGTGTCTGAGCAACTGCGAGCCCGTGCCTGCTGTTTGTGTATCACTCTCTAGG	1980	QY	3001	ATAAAGACCCCTTGCAGCTCTCGTGTCTGTGAACACTTCCCTGTGTGATTTCTGTGAGGG	3060
DB	1921	TGGGTGTCTGAGCAACTGCGAGCCCGTGCCTGCTGTTTGTGTATCACTCTCTAGG	1980	DB	3001	ATAAAGACCCCTTGCAGCTCTCGTGTCTGTGAACACTTCCCTGTGTGATTTCTGTGAGGG	3060
QY	1981	GACCTGTGTCTTCTATTTCTGTGTGACTCGTTCAATTCATCCAGGCAATTCATTGACAAAT	2040	QY	3061	GGATGTGAGAGGGGAAAGGAGGAGGAGCTGGAGAGCTGAGGCAACAGGGAGGTGAGGG	3120
DB	1981	GACCTGTGTCTTCTATTTCTGTGTGACTCGTTCAATTCATCCAGGCAATTCATTGACAAAT	2040	DB	3061	GGATGTGAGAGGGGAAAGGAGGAGGAGCTGGAGAGCTGAGGCAACAGGGAGGTGAGGG	3120
QY	2041	TATTGAGTACTTATATCTGCCAGACACACAGAGCAAAATGGTGAAGCAAGCTCACTGC	2100	QY	3121	GGACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG	3180
DB	2041	TATTGAGTACTTATATCTGCCAGACACACAGAGCAAAATGGTGAAGCAAGCTCACTGC	2100	DB	3121	GGACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG	3180
QY	2101	CCTACCTTCTGAGAGGTGACAGTTTCTCATGGAAGAGCTGCGAGAGAAATTAATAGCCA	2160	QY	3181	CAGGACCCGAGAGGCCAATGCTTTCAGGAAAGCTCAATGAACCCCAACAGCCACATTTTCT	3240
DB	2101	CCTACCTTCTGAGAGGTGACAGTTTCTCATGGAAGAGCTGCGAGAGAAATTAATAGCCA	2160	DB	3181	CAGGACCCGAGAGGCCAATGCTTTCAGGAAAGCTCAATGAACCCCAACAGCCACATTTTCT	3240
QY	2161	GCCAACTTAAACCCAGTCTGAAAGAAAGGAATAAACACCATCTTGAAGAAATTTGGCGC	2220	QY	3241	TCCCTAAGCATAGACAAATGGCAATTTGCCAATTAACAAAGAAATGTCAGAGACTTAACTGGT	3300
DB	2161	GCCAACTTAAACCCAGTCTGAAAGAAAGGAATAAACACCATCTTGAAGAAATTTGGCGC	2220	DB	3241	TCCCTAAGCATAGACAAATGGCAATTTGCCAATTAACAAAGAAATGTCAGAGACTTAACTGGT	3300
QY	2221	AGCATCCCTTAAACAGGCCACCTCCCTAGGCCCCCTGCTGCTCCATCGTGCAGGAGG	2280	QY	3301	GGTAGCTTTTGGCTGGCATTCAAAAACCTGGGCCAGAGCAAGTGGAAAAATGCCAGAGATTG	3360
DB	2221	AGCATCCCTTAAACAGGCCACCTCCCTAGGCCCCCTGCTGCTCCATCGTGCAGGAGG	2280	DB	3301	GGTAGCTTTTGGCTGGCATTCAAAAACCTGGGCCAGAGCAAGTGGAAAAATGCCAGAGATTG	3360
QY	2281	CCCCAAGCCGAGTCTTCCAGGCTCTCTCCATCAGTCAACGCGCTCAGCTGGCT	2340	QY	3361	TTAAACTTTTACCCTTGACAGCACCCACAGCAGCTCAGCAGTGAAGTGTGACAGCACGG	3420
DB	2281	CCCCAAGCCGAGTCTTCCAGGCTCTCTCCATCAGTCAACGCGCTCAGCTGGCT	2340	DB	3361	TTAAACTTTTACCCTTGACAGCACCCACAGCAGCTCAGCAGTGAAGTGTGACAGCACGG	3420
QY	2341	GCCTCGCTTCCCGTGAATCGTCTGGTGCATCTGAGCTGAGACTCTCTGGCTCCAGCT	2400	QY	3421	AGTGACCTCCAGCGAGGGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	3480
DB	2341	GCCTCGCTTCCCGTGAATCGTCTGGTGCATCTGAGCTGAGACTCTCTGGCTCCAGCT	2400	DB	3421	AGTGACCTCCAGCGAGGGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	3480
QY	2401	CCAGAAAGGAAATGGAGAGGAAACTAGTCTAAACGAGAAATCTGAGAGGAGCAAGTCTC	2460	QY	3481	ACAGATTCAATTCAGGGCAGTGGGATTCACACAGGGATTAAGTCCAGCTCATCTGG	3540
DB	2401	CCAGAAAGGAAATGGAGAGGAAACTAGTCTAAACGAGAAATCTGAGAGGAGCAAGTCTC	2460	DB	3481	ACAGATTCAATTCAGGGCAGTGGGATTCACACAGGGATTAAGTCCAGCTCATCTGG	3540
QY	2461	CTCAGAGGAAAGGGGCTTCAAGTCCAGAGAAATTCAGGAGGTGGGAGCTGAGGAG	2520	QY	3541	GTTCAGGAGGAGGGGCTATATTTGGGGGAAAAAATCAGTTTCAAGGGAAGTCCGGAGA	3600
DB	2461	CTCAGAGGAAAGGGGCTTCAAGTCCAGAGAAATTCAGGAGGTGGGAGCTGAGGAG	2520	DB	3541	GTTCAGGAGGAGGGGCTATATTTGGGGGAAAAAATCAGTTTCAAGGGAAGTCCGGAGA	3600
QY	2521	TGGGAGCTGGGCTGAGGGGGTCTGAAAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	2580	QY	3601	CCTGATTTCTAATCTATATTTTCTTTTACAGCTGAGTAATCTGAGCAAGTCAAG	3660
DB	2521	TGGGAGCTGGGCTGAGGGGGTCTGAAAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	2580	DB	3601	CCTGATTTCTAATCTATATTTTCTTTTACAGCTGAGTAATCTGAGCAAGTCAAG	3660
QY	2581	GCTGCCAGATGTTTCAAGTGTGTTTCAAGGGGCTGGGAGTTTCCGTTGCTTCTGTAGC	2640	QY	3661	GTAGTAACTGAGGCTGTAAGATTACTAGTTTCTCTCTTATAGGAACTCTTTTCTCTGT	3720
DB	2581	GCTGCCAGATGTTTCAAGTGTGTTTCAAGGGGCTGGGAGTTTCCGTTGCTTCTGTAGC	2640	DB	3661	GTAGTAACTGAGGCTGTAAGATTACTAGTTTCTCTCTTATAGGAACTCTTTTCTCTGT	3720
QY	2641	CTTTTATCTTTCTCTGTTGGAGGAGAGAGTCTATTTTCATGAAGGAGTGCAGTTTC	2700	QY	3721	GGAGTTAGCAGCAGAGGCAATCCCGTTTCTTTTAAACAGGAGAAAAATCTCTTAAGAG	3780
DB	2641	CTTTTATCTTTCTCTGTTGGAGGAGAGAGTCTATTTTCATGAAGGAGTGCAGTTTC	2700	DB	3721	GGAGTTAGCAGCAGAGGCAATCCCGTTTCTTTTAAACAGGAGAAAAATCTCTTAAGAG	3780
QY	2701	ATAAGTCACTGTTTAAATTCAGGGTGTGATGGGTTTCTTTCACAGAGGCTTTAT	2760	QY	3781	TAAAGCAACAGATTCAGGCTTAGGTCTTGTGACTATATGATGCTTTTGTGAAAAAT	3840
DB	2701	ATAAGTCACTGTTTAAATTCAGGGTGTGATGGGTTTCTTTCACAGAGGCTTTAT	2760	DB	3781	TAAAGCAACAGATTCAGGCTTAGGTCTTGTGACTATATGATGCTTTTGTGAAAAAT	3840
QY	2761	TTAATGGGAATATAGGAGGAGCTCAATTTCTTAGGGCTGTTAAATTCAGGAGAGTGC	2820	QY	3841	CATTTTCAGGAGTGTCTATCTGATTCAGAAAAATGAGACTAGTACCTTTTGTGAGCTG	3900
DB	2761	TTAATGGGAATATAGGAGGAGCTCAATTTCTTAGGGCTGTTAAATTCAGGAGAGTGC	2820	DB	3841	CATTTTCAGGAGTGTCTATCTGATTCAGAAAAATGAGACTAGTACCTTTTGTGAGCTG	3900
				QY	3901	TAAACAAACACCCAGTTGTAAATGTCTCAAGTTTCAGGCTTAACTGCAGAGAACCAATCAAAA	3960

QY 61 TCCTATTAACCTGTATAGCTCCATTCGGATGTATGTCTTTGGCAGGATGATAAGATCA 120
DB 61 TCCTATAAAGCTGTATAGCTCCATTCGGATGTATGTCTTTGGCAGGATGATAAGATCA 120
QY 121 GGAAGAAGAGATATCCACGTTAGCCAAAGTGTCCAGGCTGTCTGTCTCTTTATTTAGTGA 180
DB 121 GGAAGAAGAGATATCCACGTTAGCCAAAGTGTCCAGGCTGTCTGTCTCTTTATTTAGTGA 180
QY 181 CAGATGTTGCTCTCTGACAGAGCTATTTCTTCAGGAAACATCATCTCCATATGTTAAATC 240
DB 181 CAGATGTTGCTCTCTGACAGAGCTATTTCTTCAGGAAACATCATCTCCATATGTTAAATC 240
QY 241 CATCAACAGGAGCTTAAGAAACAGGAATGAGATGAGCTCCCAAGGAAATATGCCAG 300
DB 241 CATCAACAGGAGCTTAAGAAACAGGAATGAGATGAGCTCCCAAGGAAATATGCCAG 300
QY 301 GAGAGCAATAATATGATGAAATATAAATTTTCCCTTTGTTTAAATTTTTCAGGAAATATG 360
DB 301 GAGAGCAATAATATGATGAAATATAAATTTTCCCTTTGTTTAAATTTTTCAGGAAATATG 360
QY 361 ATGAGGACCAAAATCAATGAATTAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
DB 361 ATGAGGACCAAAATCAATGAATTAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
QY 421 TAAATTAAGTATTTGTTCTTGGGAGAGACCTCCATGTGAGCTTGTATGGGAAATGGGAA 480
DB 421 TAAATTAAGTATTTGTTCTTGGGAGAGACCTCCATGTGAGCTTGTATGGGAAATGGGAA 480
QY 481 AAACGTCAAAAGCATGATCTGATCAGATCCCAAGTGAATTAATTTTAAAAACAGAT 540
DB 481 AAACGTCAAAAGCATGATCTGATCAGATCCCAAGTGAATTAATTTTAAAAACAGAT 540
QY 541 GGCATCACTCTGGGAGGCAAGTTTCAGGAAGTCAATTTAGCAAGGACATTAACATTAAC 600
DB 541 GGCATCACTCTGGGAGGCAAGTTTCAGGAAGTCAATTTAGCAAGGACATTAACATTAAC 600
QY 601 AGCAAAATCAAAATTTCCGCAATTCAGGAGGAAATGGGACCTGGGAAAGCTTTTCAATAC 660
DB 601 AGCAAAATCAAAATTTCCGCAATTCAGGAGGAAATGGGACCTGGGAAAGCTTTTCAATAC 660
QY 661 AGTGATTAAGGCAAGTTGACCATGTTGCAACACTCCCGCTCTATACAGGGAACACAAA 720
DB 661 AGTGATTAAGGCAAGTTGACCATGTTGCAACACTCCCGCTCTATACAGGGAACACAAA 720
QY 721 ATTGACTGGGCTAAGCTGGAATTTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA 780
DB 721 ATTGACTGGGCTAAGCTGGAATTTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA 780
QY 781 GACATGTTTAAAGGCAACAGAACTTGTGAGCTTCAAGCAGCAGTGCCTCAGCA 840
DB 781 GACATGTTTAAAGGCAACAGAACTTGTGAGCTTCAAGCAGCAGTGCCTCAGCA 840
QY 841 GGGACCTCAGGCAATTTGCTTTAGGAAGCCAGTTTCTTAAGGAATCTTAAGAACTC 900
DB 841 GGGACCTCAGGCAATTTGCTTTAGGAAGCCAGTTTCTTAAGGAATCTTAAGAACTC 900
QY 901 TTGAAAGATCATGAATTTTAAACATTTTAAGTATAAAACAAATATGCGATCATATCAG 960
DB 901 TTGAAAGATCATGAATTTTAAACATTTTAAGTATAAAACAAATATGCGATCATATCAG 960
QY 961 TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAAAGGATACGTTGCCAGTCC 1020
DB 961 TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAAAGGATACGTTGCCAGTCC 1020
QY 1021 GGATAGGTCAGAAATCATTAGAAATCATGTGTCCCACTCTTAACATTTTTCAGAAATGATC 1080
DB 1021 GGATAGGTCAGAAATCATTAGAAATCATGTGTCCCACTCTTAACATTTTTCAGAAATGATC 1080
QY 1081 TGTATAGCCCTCACAACAGGCCCGATGTGTCTGACCTTCAACACATCTTCAACCCAA 1140
DB 1081 TGTATAGCCCTCACAACAGGCCCGATGTGTCTGACCTTCAACACATCTTCAACCCAA 1140
QY 1141 GTGCCTCAACCATTTGTTAAGTGTCTATCTCAGTAGGTCCCATTTACAAATGCCACCTCCCC 1200

DB 1141 GTGCCTCAACCATTTGTTAAGTGTCTATCTCAGTAGGTCCCATTTACAAATGCCACCTCCCC 1200
QY 1201 TGTGAGGCCCATTCGCGCTCCACAGGAAGTCTCCCACTCTAGACTTTGTCATCAGATGT 1260
DB 1201 TGTGAGGCCCATTCGCGCTCCACAGGAAGTCTCCCACTCTAGACTTTGTCATCAGATGT 1260
QY 1261 TACAGCCAGAAGCTCCGCTGAGGCTGAGGCTCTGTGTCTTACACCTTACCTGTATGTCTAC 1320
DB 1261 TACAGCCAGAAGCTCCGCTGAGGCTGAGGCTCTGTGTCTTACACCTTACCTGTATGTCTAC 1320
QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCCTCC 1380
DB 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCCTCC 1380
QY 1381 CGCTGAGTGGGACTACAGGCGCACGCGCGCTAATTTTGTATTTAGTAGAGTGG 1440
DB 1381 CGCTGAGTGGGACTACAGGCGCACGCGCGCTAATTTTGTATTTAGTAGAGTGG 1440
QY 1441 GTTTCACCATATTAAGCCGCTGCTTGAATCTCTGACCTCAGGTGATCCACCCACCTC 1500
DB 1441 GTTTCACCATATTAAGCCGCTGCTTGAATCTCTGACCTCAGGTGATCCACCCACCTC 1500
QY 1501 AGCCTCTTAAAGTGTCTGGGATTTACAGGATGAGTCACCGCCCGCCGCAAGGTTCA 1560
DB 1501 AGCCTCTTAAAGTGTCTGGGATTTACAGGATGAGTCACCGCCCGCCGCAAGGTTCA 1560
QY 1561 TTAATAAGGAATACTTGAATGTTTACTTAAACCAAGGGAACAGACAAAGCTGGA 1620
DB 1561 TTAATAAGGAATACTTGAATGTTTACTTAAACCAAGGGAACAGACAAAGCTGGA 1620
QY 1621 TAAATTCAGGATTTCTGGGATGGGAATGTTGCCATGAGCTGCTGCTAGTCCAGAC 1680
DB 1621 TAAATTCAGGATTTCTGGGATGGGAATGTTGCCATGAGCTGCTGCTAGTCCAGAC 1680
QY 1681 CACTGGTCTCATCACTTTCTTCCCTCATCTCTCAATTTTCAAGCTAAGTTTACATTTTAT 1740
DB 1681 CACTGGTCTCATCACTTTCTTCCCTCATCTCTCAATTTTCAAGCTAAGTTTACATTTTAT 1740
QY 1741 CACATGTTTTGTGTTAAGCTTCCACATCTGTTACTGAAATAGATATACATAACTAG 1800
DB 1741 CACATGTTTTGTGTTAAGCTTCCACATCTGTTACTGAAATAGATATACATAACTAG 1800
QY 1801 TTCCATTTTGGGCTCATCTGTGTGTATAGGAGGAGGCAATACCCAGAGACTCTCT 1860
DB 1801 TTCCATTTTGGGCTCATCTGTGTGTATAGGAGGAGGCAATACCCAGAGACTCTCT 1860
QY 1861 TGAAGCCCCCGGAGAGGTTTCTCTCCAGCTGGGGAGCCCTGCAAGCACCCGGGTCC 1920
DB 1861 TGAAGCCCCCGGAGAGGTTTCTCTCCAGCTGGGGAGCCCTGCAAGCACCCGGGTCC 1920
QY 1921 TGGGTGCTCTGAGCAACCTGCGCCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1980
DB 1921 TGGGTGCTCTGAGCAACCTGCGCCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1980
QY 1981 GACCTGTTGCTTTCTTATTTCTGTGACTCTGTTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2040
DB 1981 GACCTGTTGCTTTCTTATTTCTGTGACTCTGTTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2040
QY 2041 TATTGAGTACTTATATCTGCGAGACACAGAGACAAAATGTTAGCAAAAGCAGTCACTGC 2100
DB 2041 TATTGAGTACTTATATCTGCGAGACACAGAGACAAAATGTTAGCAAAAGCAGTCACTGC 2100
QY 2101 CTTACCTTCTGAGGAGTGAAGTTTCTCATGGAAGACGTCGAGAGAAAATTAATAGCCA 2160
DB 2101 CTTACCTTCTGAGGAGTGAAGTTTCTCATGGAAGACGTCGAGAGAAAATTAATAGCCA 2160
QY 2161 GCCAATTTAAACCCAGTCTGAAAGAAAGGAAATAAACCACTTTTGAAGAAATTTGTCGC 2220
DB 2161 GCCAATTTAAACCCAGTCTGAAAGAAAGGAAATAAACCACTTTTGAAGAAATTTGTCGC 2220
QY 2221 AGCATCCCTTAAACAGGCCACCTCCCTAGCGCCCCCTGCTGCTCCTCATCTGTCGCCGAGG 2280

Db 2221 AGCATCCCTTAACAAAGGCGACCTCCCTAGCGCCCTCTGCTGCTCCATGCTGCTCCCGGAGG 2280
Qy 2281 CCCCAAGCCGAGTCTTCCAAAGCCTCTCTCCATCAGTCAAGGCTGAGCTGAGCT 2340
Db 2281 CCCCAAGCCGAGTCTTCCAAAGCCTCTCTCCATCAGTCAAGGCTGAGCTGAGCT 2340
Qy 2341 GCCTCGCTTCCCGTGAATCGTCTCGTGTGATCTGAGCTGGAGCTCTCTGGCTCCAGCT 2400
Db 2341 GCCTCGCTTCCCGTGAATCGTCTCGTGTGATCTGAGCTGGAGCTCTCTGGCTCCAGCT 2400
Qy 2401 CCAGAAAGAAATGAGAGGGGAACTAGTCTAAACGAGAACTGAGAGGGGAGAGTCTTC 2460
Db 2401 CCAGAAAGAAATGAGAGGGGAACTAGTCTAAACGAGAACTGAGAGGGGAGAGTCTTC 2460
Qy 2461 CTCAGAGGAAGGGGCTCCACGCTCCAGAGAACTCCAGAGGTGGGAGCTGCAAGGAG 2520
Db 2461 CTCAGAGGAAGGGGCTCCACGCTCCAGAGAACTCCAGAGGTGGGAGCTGCAAGGAG 2520
Qy 2521 TGGGACGCTGGGCTGAGCGGGTCTGAAAGGCGAGGAGGTGAAAGGGCAAGGCTGAA 2580
Db 2521 TGGGACGCTGGGCTGAGCGGGTCTGAAAGGCGAGGAGGTGAAAGGGCAAGGCTGAA 2580
Qy 2581 GCTGCCAGATGTTCAAGTGTGTTTCAAGGGCTGGGAGTTCCTGCTCTCTGTGAGC 2640
Db 2581 GCTGCCAGATGTTCAAGTGTGTTTCAAGGGCTGGGAGTTCCTGCTCTCTGTGAGC 2640
Qy 2641 CTTTTTATCTTTCTCTGCTGGAGGAGAGTCTATTTTTCATGAAGGGATGAGTTC 2700
Db 2641 CTTTTTATCTTTCTCTGCTGGAGGAGAGTCTATTTTTCATGAAGGGATGAGTTC 2700
Qy 2701 ATAAAGTCAGCTGTTTAAATTCAGGGTGTGATGGTTCCTTTCAGAAAGGCTTAT 2760
Db 2701 ATAAAGTCAGCTGTTTAAATTCAGGGTGTGATGGTTCCTTTCAGAAAGGCTTAT 2760
Qy 2761 TTAATGGGAATATAGGAAGGAGCTCATTTCTAGCGGTAAATTCAGGAAGAGTGAC 2820
Db 2761 TTAATGGGAATATAGGAAGGAGCTCATTTCTAGCGGTAAATTCAGGAAGAGTGAC 2820
Qy 2821 TGGAGTCTTTTCTTCTGCTCTGCGCACTACTCAGCCCTGCTGGTGGACTGGCTTA 2880
Db 2821 TGGAGTCTTTTCTTCTGCTCTGCGCACTACTCAGCCCTGCTGGTGGACTGGCTTA 2880
Qy 2881 TGCAGAGCGTTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTCTGCTTCGCCATT 2940
Db 2881 TGCAGAGCGTTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTCTGCTTCGCCATT 2940
Qy 2941 GGTGGCTGTGCGACCGTGGGCAAGTGTCTCTCTCCCTGGGCCATAGTCTTCTCTGCT 3000
Db 2941 GGTGGCTGTGCGACCGTGGGCAAGTGTCTCTCTCCCTGGGCCATAGTCTTCTCTGCT 3000
Qy 3001 ATAAAGCCCTTGCAGCTCTGCTGTCTGTGAACTCTCCCTGTGATCTCTGTGAGGG 3060
Db 3001 ATAAAGCCCTTGCAGCTCTGCTGTCTGTGAACTCTCCCTGTGATCTCTGTGAGGG 3060
Qy 3061 GGATGTTGAGAGGGGAGGAGGAGCTGAGAGCTGAGCCACAGGGAGGTGGAGGG 3120
Db 3061 GGATGTTGAGAGGGGAGGAGGAGCTGAGAGCTGAGCCACAGGGAGGTGGAGGG 3120
Qy 3121 GGACAGGAAGGAGGAGGAGCTGGGTCTCCATCAGTCTCTCAGTATCAGCTCAGACTC 3180
Db 3121 GGACAGGAAGGAGGAGGAGCTGGGTCTCCATCAGTCTCTCAGTATCAGCTCAGACTC 3180
Qy 3181 CAGGACCGAGGAGCCCAATGCTTCCAGGAAGCTCAATGAACCCCAAGCCACATTTTCCT 3240
Db 3181 CAGGACCGAGGAGCCCAATGCTTCCAGGAAGCTCAATGAACCCCAAGCCACATTTTCCT 3240
Qy 3241 TCCCTAAGCATAGACATGGCATTTGCAATTAACCAAAAGAAATGCGAGACTAATCGGT 3300
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Qy 3301 GGTAAGCTTTTCTGCGCATTCAAAACTGGGCCAGAGCAAGTGGAAATGCCAGAGATTG 3360
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Qy 3361 TTAACTTTTCCACCTGACGACGACCCACGAGCTCAGCTGACTGCTGACGACGCG 3420
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Db 3960 AAGAAATAGAATCTTTAGAGCAAACTGTGTTTCTCCAC - TCTGGAGGTGAGTCTGCCAGGG 4019
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Qy 4200 GGGATTTAACTTACAGTCCAGAAAGCCTGTGAATTTGAATGAGGAAAGAAATTTACATTT 4259
Db 4200 GGGATTTAACTTACAGTCCAGAAAGCCTGTGAATTTGAATGAGGAAAGAAATTTACATTT 4259
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Db 4380 TTTTATCTATATTTACAGTCTTGGAGATCGTTGTGAAGTGAATATTTTATCTCAAACT 4439

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Db	1621	TAAATTCAGGATTCCTTGGGATGGGGAATGGTCCCATGAGTCCCTGCTAGTCCAGAC	1680
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Db	1681	CACGTGCTCATCACTTTCTCCCTCATCTCACTCACTTTCAGGCTAAGTTACCAATTTAT	1740
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Db	1861	TGAAGCCCCCGCAGAGGTTTCTCTCCAGCTGGGGAGCCCTGCAAGCACCCGGGCTC	1920
Qy	1921	TGGGTGCTGAGCAACTGCTCCAGAGCCGCTGCCACTGGTTTGTATCACTCTCAGG	1980
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Db	2041	TATTGAGTACTTATATCTGCCAGACACAGAGACAAATGGTGAGCAAGCACTGC	2100
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Db	2101	CCTACCTTCTGAGGTGACAGTTTCTCATGGAAGAGCTGCAGAGAAATTAATAGCA	2160
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Qy	2221	AGCATCCCTTAAACAGGCCACCTCCTAGCGCCCTGCTGCTCCATCTGCGCGGAGG	2280
Db	2221	AGCATCCCTTAAACAGGCCACCTCCTAGCGCCCTGCTGCTCCATCTGCGCGGAGG	2280
Qy	2281	CCCCAAGCCGAGTCTTCCAGGCTCCTCCTCATCAGTCAAGCGCTGCAGCTGCGCT	2340
Db	2281	CCCCAAGCCGAGTCTTCCAGGCTCCTCCTCATCAGTCAAGCGCTGCAGCTGCGCT	2340
Qy	2341	GCCTCGCTTCCGCTGAATCGTCTGTGTCATCTGAGCTGAGACTCCTTGGCTCCAGGCT	2400
Db	2341	GCCTCGCTTCCGCTGAATCGTCTGTGTCATCTGAGCTGAGACTCCTTGGCTCCAGGCT	2400
Qy	2401	CCAGAAAGGAAATGGAGAGGAAACTAGTCTAACCGGAGAAATCTGGAGGGGACAGTGTTC	2460
Db	2401	CCAGAAAGGAAATGGAGAGGAAACTAGTCTAACCGGAGAAATCTGGAGGGGACAGTGTTC	2460
Qy	2461	CTCAGAGGAAAGGGGCTTCCAGTCCAGAGAAATTCAGAGGTGGGGACTGCGGGAG	2520
Db	2461	CTCAGAGGAAAGGGGCTTCCAGTCCAGAGAAATTCAGAGGTGGGGACTGCGGGAG	2520
Qy	2521	TGGGACGCTGGGCTCAGCGGGTGTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	2580
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Qy	2581	GCTGCCAGATGTTTCAGTGTGTTTTCACGGGGCTGGAGTGTTCCTGCTTCTCTGAGC	2640
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Qy	2761	TTAATGGGAATATAGGAAGGAGCTCATTTCTAGCCGTTAATTCACGAAGGATGAC	2820
Db	2761	TTAATGGGAATATAGGAAGGAGCTCATTTCTAGCCGTTAATTCACGAAGGATGAC	2820
Qy	2821	TGGAGTCTTTTCTTTCATGCTCTTCTGGGCAACTACTCAGCCCTGTGGTGGACTTGA	2880
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Qy	2881	TGCAAGCGGTGCGAAGGAGCTGCGGATCAGGAGACTCGGTTTCTTCTGGTCTGCAAT	2940
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Qy	2941	GGTTGGCTGTGCGACCGTGGGCAAGTGTCTCTTCCCTGGGCGCATAGTCTCTGCT	3000
Db	2941	GGTTGGCTGTGCGACCGTGGGCAAGTGTCTCTTCCCTGGGCGCATAGTCTCTGCT	3000
Qy	3001	ATAAAGACCTTTCAGTCTCTGTTCTGTGAACACTTCCCTGTGATTTCTCTGTGAGGG	3060
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Db	3061	GGATGTTGAGGGGAAAGGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGGAGGAGG	3120
Qy	3121	GGACGAGGAGGAGGAGGAGGAGCTGGGTGCTCATCAGTCTCCTCATCAGTCACTC	3180
Db	3121	GGACGAGGAGGAGGAGGAGGAGCTGGGTGCTCATCAGTCTCCTCATCAGTCACTC	3180
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Db	3241	TCCCTAAGCATAGACAAATGGCATTTGCCAATTAACCAAAAAGAAATGAGAGACTTAACTGGT	3300
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[illegible]

RESULT 8

US-09-056-285A-1
Sequence 1, Application US/09056285A
Patent No. 6403307
GENERAL INFORMATION:
APPLICANT: Stone, Edwin M.
Sheffield, Val C.
Alward, Wallace L.M.
Fingert, John
TITLE OF INVENTION: GLAUCOMA THERAPEUTICS AND DIAGNOSTICS
NUMBER OF SEQUENCES: 43
CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY, HOAG & ELIOT LLP
STREET: One Post Office Square
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02109-2170
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent in Release #1.0. Version #1.30

GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 15852
; LENGTH: 205044
; TYPE: DNA
; ORGANISM: Human
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)...(205044)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-15852

Query Match 3.5%; Score 184.6; DB 3; Length 205044;
Best Local Similarity 78.0%; Pred. No. 3.6e-34;
Matches 238; Conservative 0; Mismatches 59; Indels 8; Gaps 1;

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Search completed: January 26, 2006, 04:23:24
Job time : 614 secs

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OM nucleic - nucleic search, using sw model

Run on: January 26, 2006, 03:55:20 ; Search time 2551 Seconds
(without alignments)
17086.590 Million cell updates/sec

Title: US-09-227-881-34

Perfect score: 5271

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Total number of hits satisfying chosen parameters: 19587084

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

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9: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq.*
10: /cgn2_6/ptodata/1/pubpna/US11_PUBCOMB.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	5271	100.0	5271	5	US-10-244-633-34
2	5271	100.0	6169	5	US-10-244-633-3
3	5246.4	99.5	5300	3	US-09-985-637A-1
4	5246.4	99.5	5300	5	US-10-244-633-1
5	5246.4	99.5	5300	7	US-10-741-339-1
6	5232.4	99.3	37252	5	US-10-087-192-1228
7	5224.4	99.1	5304	5	US-10-244-633-2
8	4017.4	76.2	6000	9	US-10-509-595-1
9	1804.4	34.2	2800	3	US-09-952-464A-1
10	1804.4	34.2	2800	6	US-10-017-870-10
11	1804.4	34.2	2800	8	US-10-278-698-294
12	1804.4	34.2	2800	8	US-10-278-698-808
13	1804.4	34.2	2800	9	US-10-803-557-10
14	1804.4	34.2	2800	9	US-10-956-243-1
15	631.6	12.0	632	4	US-09-925-065A-905501
16	629.6	11.9	630	4	US-09-925-065A-917000
17	546	10.4	571	4	US-09-925-065A-346049
18	394.6	7.5	545	4	US-09-925-065A-346051
19	394.4	7.5	1086	7	US-10-240-425-1586
20	394.4	7.5	1086	8	US-10-723-860-927
21	394.4	7.5	1086	9	US-10-756-149-932
22	393	7.5	545	4	US-09-925-065A-346050
23	284	5.4	296	8	US-10-674-124A-1442

C	24	284	5.4	296	8	US-10-674-124A-1443	Sequence 1443, Ap
	25	227	4.3	227	5	US-10-244-633-38	Sequence 38, Appl
	26	227	4.3	283	5	US-10-244-633-37	Sequence 37, Appl
	27	222.6	4.2	30057	5	US-10-087-192-1225	Sequence 1225, Ap
C	28	194.6	3.7	337344	8	US-10-388-838-58	Sequence 58, Appl
C	29	189.8	3.6	37443	8	US-10-719-993-6853	Sequence 6853, Ap
C	30	185.4	3.5	86361	7	US-10-741-601-5702	Sequence 5702, Ap
C	31	185.4	3.5	86361	8	US-10-741-600-17803	Sequence 17803, A
C	32	184.4	3.5	1744	6	US-10-104-047-1845	Sequence 1845, Ap
C	33	184.2	3.5	70779	5	US-10-087-192-1012	Sequence 1012, Ap
C	34	183.2	3.5	41936	3	US-09-967-768A-116	Sequence 116, App
C	35	183.2	3.5	41936	9	US-10-843-641A-6261	Sequence 6261, Ap
C	36	183.2	3.5	104245	6	US-10-160-807-4	Sequence 4, Appli
C	37	183.2	3.5	104245	7	US-10-655-847-4	Sequence 4, Appli
C	38	183.2	3.5	170245	7	US-10-717-597-322	Sequence 322, App
C	39	182.2	3.5	5367	9	US-10-450-763-16437	Sequence 16437, A
C	40	181.2	3.4	560	4	US-09-925-065A-127303	Sequence 127303,
C	41	181.2	3.4	196063	7	US-10-322-281-612	Sequence 612, App
C	42	180.6	3.4	1620	4	US-09-925-065A-67041	Sequence 67041, A
C	43	180.6	3.4	1620	4	US-09-925-065A-67042	Sequence 67042, A
C	44	180.6	3.4	39725	6	US-10-017-161-1611	Sequence 1611, Ap
C	45	180.6	3.4	39725	6	US-10-292-798-1285	Sequence 1285, Ap

ALIGNMENTS

RESULT 1

US-10-244-633-34
; Sequence 34, Application US/10244633
; Publication No. US20030068640A1
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis, Prognosis And Treatment Of Glaucoma And Related Disorders
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: 07425.0057.US01
; CURRENT APPLICATION NUMBER: US/10/244,633
; PRIOR FILING DATE: 2002-09-17
; PRIOR APPLICATION NUMBER: US/09/306,828
; PRIOR FILING DATE: 1999-05-07
; PRIOR APPLICATION NUMBER: US 09/227,881
; PRIOR FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 34
; LENGTH: 5271
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-244-633-34

Query Match 100.0%; Score 5271; DB 5; Length 5271;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 5271; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAATGAAATGAGATAACCAATGTGAAAG	60
Db	1	ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAATGAAATGAGATAACCAATGTGAAAG	60
Qy	61	TCCTATAAATGTATAGCTCCATTCGGATGTATGTCTTTGGCAGGATGATAAAGATCA	120
Db	61	TCCTATAAATGTATAGCTCCATTCGGATGTATGTCTTTGGCAGGATGATAAAGATCA	120
Qy	121	GGAGAAGAGATATCCACGTTAGCCAAAGTGTCCAGGCTGTCTGCTCTTTATTTTATGTA	180
Db	121	GGAGAAGAGATATCCACGTTAGCCAAAGTGTCCAGGCTGTCTGCTCTTTATTTTATGTA	180
Qy	181	CAGATGTTCTCTGTGACAGAAGCTATTTCTTCAGGAAACATCATCCATATGTTAAATC	240
Db	181	CAGATGTTCTCTGTGACAGAAGCTATTTCTTCAGGAAACATCATCCATATGTTAAATC	240

QY	241	CATCAACAGGAGCTAAGAAACAGGAATGAGATGGGACCTTGCCCAAGGAAAAATGCCAG	300
DB	241	CATCAACAGGAGCTAAGAAACAGGAATGAGATGGGACCTTGCCCAAGGAAAAATGCCAG	300
QY	301	GAGAGCAAAATGAATGAATAAATACTTTTCCCTTTGTTTTTAAATTTAGGAAAAATG	360
DB	301	GAGAGCAAAATGAATGAATAAATACTTTTCCCTTTGTTTTTAAATTTAGGAAAAATG	360
QY	361	ATCAGAGCAAAATCAATGAATAAGGAAAAACAGCTCAGAAAAAAGATGTTTCCAAATGG	420
DB	361	ATCAGAGCAAAATCAATGAATAAGGAAAAACAGCTCAGAAAAAAGATGTTTCCAAATGG	420
QY	421	TAATTAAGTATTTGTTCTTTGGGAAGAGACCTCCATGTGAGCTTGATGGAAAAATGGAA	480
DB	421	TAATTAAGTATTTGTTCTTTGGGAAGAGACCTCCATGTGAGCTTGATGGAAAAATGGAA	480
QY	481	AAACGTCAAAAGCATGATCTGATCAGATCCCAAAGTGGATTAATTTTAAAAAACAGAT	540
DB	481	AAACGTCAAAAGCATGATCTGATCAGATCCCAAAGTGGATTAATTTTAAAAAACAGAT	540
QY	541	GGCATCACTCTGGGGAGCAAGTTTCAGAAAGGTCATGTTAGCAAAAGGACATAACAATAAC	600
DB	541	GGCATCACTCTGGGGAGCAAGTTTCAGAAAGGTCATGTTAGCAAAAGGACATAACAATAAC	600
QY	601	AGCAAAATCAAAATTTCCGCAATTCAGGAGGAAAAATGGGACCTGGGAAAGCTTTCAATAAC	660
DB	601	AGCAAAATCAAAATTTCCGCAATTCAGGAGGAAAAATGGGACCTGGGAAAGCTTTCAATAAC	660
QY	661	AGTGAATTAGGAGTTGACATGTTTCGCAACACCTCCCGCTATACCAAGGGAACACAAA	720
DB	661	AGTGAATTAGGAGTTGACATGTTTCGCAACACCTCCCGCTATACCAAGGGAACACAAA	720
QY	721	ATTGACTGGGCTAAGCCTGACATTTCAAGGGAATATGAAAACTGAGAGCAAAACAAA	780
DB	721	ATTGACTGGGCTAAGCCTGACATTTCAAGGGAATATGAAAACTGAGAGCAAAACAAA	780
QY	781	GACATGGTTAAAGGCAACAGAAACATTTGTGAGCCTTCAAAGCAGCAGTGCCCTCAGCA	840
DB	781	GACATGGTTAAAGGCAACAGAAACATTTGTGAGCCTTCAAAGCAGCAGTGCCCTCAGCA	840
QY	841	GGGACCTTGAGGCAATTTGCTTTAGGAGGCGCAGTTTCTTAAGGAATCTTAAGAAATC	900
DB	841	GGGACCTTGAGGCAATTTGCTTTAGGAGGCGCAGTTTCTTAAGGAATCTTAAGAAATC	900
QY	901	TTGAAAGCATCATGAATTTTAAACATTTTAAAGTATAAAACAAATATGCGATGCAATACAG	960
DB	901	TTGAAAGCATCATGAATTTTAAACATTTTAAAGTATAAAACAAATATGCGATGCAATACAG	960
QY	961	TTTAGACATGGGTCCTCAATTTTATAAAGTCAGGCAATCAAAGGATAACGTCCTCCAGCTCC	1020
DB	961	TTTAGACATGGGTCCTCAATTTTATAAAGTCAGGCAATCAAAGGATAACGTCCTCCAGCTCC	1020
QY	1021	GGATAGTCAGAAATCATTAAGAAATCATGTGTCTCCCATCTTAATCTTTTTCAGNATGATC	1080
DB	1021	GGATAGTCAGAAATCATTAAGAAATCATGTGTCTCCCATCTTAATCTTTTTCAGNATGATC	1080
QY	1081	TGTCATAGCCTCACACACAGGCGCGATGTGTCTGACCTACAACACATCTACAACCCAA	1140
DB	1081	TGTCATAGCCTCACACACAGGCGCGATGTGTCTGACCTACAACACATCTACAACCCAA	1140
QY	1141	GTGCTTCAACCATTTGTTAAAGTGTGCATCTCAGTAGGTCCCATTAACAAATGCACTCCCTCC	1200
DB	1141	GTGCTTCAACCATTTGTTAAAGTGTGCATCTCAGTAGGTCCCATTAACAAATGCACTCCCTCC	1200
QY	1201	TGTCAGGCGCATCCCGCTCCACAGGAATCTCCCGCATCTAGACTTCTGCATCAGCATGT	1260
DB	1201	TGTCAGGCGCATCCCGCTCCACAGGAATCTCCCGCATCTAGACTTCTGCATCAGCATGT	1260
QY	1261	TACAGCCAGAGCTCCCGTAGGGGTGAGGGTCTGTGTCTTACACCTACTGTATGCTCTAC	1320
DB	1261	TACAGCCAGAGCTCCCGTAGGGGTGAGGGTCTGTGTCTTACACCTACTGTATGCTCTAC	1320
QY	1321	ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC	1380
DB	1321	ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC	1380
QY	1381	CCGCTAGCTGGGACTACAGGCGCAGCCCGGCTAAATTTTGTATTTGTAGTAGAGATGG	1440
DB	1381	CCGCTAGCTGGGACTACAGGCGCAGCCCGGCTAAATTTTGTATTTGTAGTAGAGATGG	1440
QY	1441	GTTTCCCATATTTAGCCCGGCTTGCTTGAACCTCCTGACCTCAGTGATCCACCCACCTC	1500
DB	1441	GTTTCCCATATTTAGCCCGGCTTGCTTGAACCTCCTGACCTCAGTGATCCACCCACCTC	1500
QY	1501	AGCCTCTTAAAGTGTGGGATTTACAGGCATGAGTCAACCGCGCCGCGCAAGGGTCAAGTGT	1560
DB	1501	AGCCTCTTAAAGTGTGGGATTTACAGGCATGAGTCAACCGCGCCGCGCAAGGGTCAAGTGT	1560
QY	1561	TTAATAAGGAATAAATTTGAATGTTTACTTAAACCAACAGGGAACAGACAAAAAGCTGTGA	1620
DB	1561	TTAATAAGGAATAAATTTGAATGTTTACTTAAACCAACAGGGAACAGACAAAAAGCTGTGA	1620
QY	1621	TAAATTCAGGGAATCTTTGGGATGGGAAATGGTGCATGAGCTGCCTGCTAGTCCCGAC	1680
DB	1621	TAAATTCAGGGAATCTTTGGGATGGGAAATGGTGCATGAGCTGCCTGCTAGTCCCGAC	1680
QY	1681	CACTGGTCTCATCACTTTTCTCCCTCATCTCTTCCAGGCTAAGTTTACCAATTTTATT	1740
DB	1681	CACTGGTCTCATCACTTTTCTCCCTCATCTCTTCCAGGCTAAGTTTACCAATTTTATT	1740
QY	1741	CACCATGCTTTTGTGTGAAGCCTTCCACATCGTTTACTGAAATTAAGAGTATACATAAAGTAG	1800
DB	1741	CACCATGCTTTTGTGTGAAGCCTTCCACATCGTTTACTGAAATTAAGAGTATACATAAAGTAG	1800
QY	1801	TTCCATTTGGGGCAATCTGTGTGTGTATAGGGAGGAGGATACCCAGAGACTCCT	1860
DB	1801	TTCCATTTGGGGCAATCTGTGTGTGTATAGGGAGGAGGATACCCAGAGACTCCT	1860
QY	1861	TGAAGCCCGCGGAGAGGTTTCTCTCCAGCTGGGGGAGCCCTGCAAGCACCCCGGGTCC	1920
DB	1861	TGAAGCCCGCGGAGAGGTTTCTCTCCAGCTGGGGGAGCCCTGCAAGCACCCCGGGTCC	1920
QY	1921	TGGGTGTCTGAGCAACCTGCCAGCCCGTGCCACTGCTGTTTGTATCACTCTCTAGG	1980
DB	1921	TGGGTGTCTGAGCAACCTGCCAGCCCGTGCCACTGCTGTTTGTATCACTCTCTAGG	1980
QY	1981	GACCTGTTGCTTTCTATTTCTGTGTGACTCGTTTCAATTCAGGCAATTCATTTGACAAAT	2040
DB	1981	GACCTGTTGCTTTCTATTTCTGTGTGACTCGTTTCAATTCAGGCAATTCATTTGACAAAT	2040
QY	2041	TATTGAGTACTTATATCTGCCAGACACCAAGACAAATGGTGAGCAAAAGCAGTCACTGC	2100
DB	2041	TATTGAGTACTTATATCTGCCAGACACCAAGACAAATGGTGAGCAAAAGCAGTCACTGC	2100
QY	2101	CCTACCTCTGTTGAGGTGACAGTTTCTCATGGAAGAGAGTGCAGAGAAAAATTAATAGCCA	2160
DB	2101	CCTACCTCTGTTGAGGTGACAGTTTCTCATGGAAGAGAGTGCAGAGAAAAATTAATAGCCA	2160
QY	2161	GCCAACTTAAACCCAGTGTGAAGAAATAAACCACTCTTTGAAGAAATTTGTGGC	2220
DB	2161	GCCAACTTAAACCCAGTGTGAAGAAATAAACCACTCTTTGAAGAAATTTGTGGC	2220
QY	2221	AGCATCCCTTAAACAGGCCACCTCCCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGG	2280
DB	2221	AGCATCCCTTAAACAGGCCACCTCCCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGG	2280
QY	2281	CCCCCAAGCCCGAGTCTTCCAGCCCTCTCTCCATCAGTCAAGGGCTGAGGTGGCCT	2340
DB	2281	CCCCCAAGCCCGAGTCTTCCAGCCCTCTCTCCATCAGTCAAGGGCTGAGGTGGCCT	2340
QY	2341	GCCTCGCTTCCCGTGAATCGTCTGTCATCTGAGCTGGAGACTCTTGGCTCCAGGCT	2400
DB	2341	GCCTCGCTTCCCGTGAATCGTCTGTCATCTGAGCTGGAGACTCTTGGCTCCAGGCT	2400
QY	2401	CCAGAAAGGAATGGAGAGGGAATAGTCTAAGGAGAACTCTGGAGGGGACAGTGTTC	2460

|||||
2401 CCAGAAAGGAAATCGAGGGGAACTAGTCTAA CGGAGAACTGGAGGGGACAGTGTTC 2460
Qy
2461 CTCAGAGGAAAGGGGCTCCAGCTCCAGGAGAAATTCAGAGAGTGGGGA CTG CAGGGAG 2520
Db
2461 CTCAGAGGAAAGGGGCTCCAGCTCCAGGAGAAATTCAGAGAGTGGGGA CTG CAGGGAG 2520
Qy
2521 TGGGGACGCTGGGCTCGAGCGGGTCTGAAAGGCGAGGAAGGTGAAAGGGGCAAGGCTGAA 2580
Db
2521 TGGGGACGCTGGGCTCGAGCGGGTCTGAAAGGCGAGGAAGGTGAAAGGGGCAAGGCTGAA 2580
Qy
2581 GCTGCCAGATGTTCACTGTTGTTTTCAGGGGCTGGGAGTTTTCCGTTGCTTCTCTGTGAGC 2640
Db
2581 GCTGCCAGATGTTCACTGTTGTTTTCAGGGGCTGGGAGTTTTCCGTTGCTTCTCTGTGAGC 2640
Qy
2641 CTTTTTATCTTTCTCTGCTTGGAGGAGAAAGTCTATTTTCATGAAGGGATG CAGTGTTC 2700
Db
2641 CTTTTTATCTTTCTCTGCTTGGAGGAGAAAGTCTATTTTCATGAAGGGATG CAGTGTTC 2700
Qy
2701 ATAAAGTCAGCTGTTAAAAATTCAGGGTGTGCATGGGTTTTTCTTCACGAAGGCTTTAT 2760
Db
2701 ATAAAGTCAGCTGTTAAAAATTCAGGGTGTGCATGGGTTTTTCTTCACGAAGGCTTTAT 2760
Qy
2761 TTAATGGGAATATAGGAAGCGAGCTCATTTCTAGGCGGTTAAATTCACGGAAGAGTGAC 2820
Db
2761 TTAATGGGAATATAGGAAGCGAGCTCATTTCTAGGCGGTTAAATTCACGGAAGAGTGAC 2820
Qy
2821 TGGAGTCTTTTCTTTCATGCTCTCTGGGCAACTACTCAGGCCCTGTGGTGGACTTGGCTTA 2880
Db
2821 TGGAGTCTTTTCTTTCATGCTCTCTGGGCAACTACTCAGGCCCTGTGGTGGACTTGGCTTA 2880
Qy
2881 TGCAGACGCTCGAAACCTTGGAAATCAGGAGACTCGGTTTTCTTCTGCTTCTGCCATT 2940
Db
2881 TGCAGACGCTCGAAACCTTGGAAATCAGGAGACTCGGTTTTCTTCTGCTTCTGCCATT 2940
Qy
2941 GGTGGCTGTGCGACCTGGGCAAGTGTCTCTCTCCCTGGGCAATAGTCTTCTCTGCT 3000
Db
2941 GGTGGCTGTGCGACCTGGGCAAGTGTCTCTCTCCCTGGGCAATAGTCTTCTCTGCT 3000
Qy
3001 ATAAAGACCTTGCAGCTCTGTTGTTCTGTGAACACTTCCCTGTGATTTCTCTGTGAGGG 3060
Db
3001 ATAAAGACCTTGCAGCTCTGTTGTTCTGTGAACACTTCCCTGTGATTTCTCTGTGAGGG 3060
Qy
3061 GGAATGTTGAGGGGAGGAGGAGCTGGAGCAGCTGAGCAGCTGAGCCAGGGAGGTGGAGG 3120
Db
3061 GGAATGTTGAGGGGAGGAGGAGCTGGAGCAGCTGAGCAGCTGAGCCAGGGAGGTGGAGG 3120
Qy
3121 GGACAGGAAGGCGAGGAGAGCTGGGTGCTCCATCAGTCCCTCAGTCACTGATCAGCTCAGACTC 3180
Db
3121 GGACAGGAAGGCGAGGAGAGCTGGGTGCTCCATCAGTCCCTCAGTCACTGATCAGCTCAGACTC 3180
Qy
3181 CAGGACCGAGGCGCAATGCTTCAGGAAAGCTCAATGAACCCAAACAGCCACATTTTCT 3240
Db
3181 CAGGACCGAGGCGCAATGCTTCAGGAAAGCTCAATGAACCCAAACAGCCACATTTTCT 3240
Qy
3241 TCCCTAAGCATAGACATGCAATTTGCCAATAACCCAAAAGAGATGAGAGACTAATCTGTT 3300
Db
3241 TCCCTAAGCATAGACATGCAATTTGCCAATAACCCAAAAGAGATGAGAGACTAATCTGTT 3300
Qy
3301 GGTAGCTTTTCTGCTGCAATCAAAAACTGGGCGAGAGCAAGTGGAAAAATGCCAGAGATTG 3360
Db
3301 GGTAGCTTTTCTGCTGCAATCAAAAACTGGGCGAGAGCAAGTGGAAAAATGCCAGAGATTG 3360
Qy
3361 TTAACCTTTTCACTGAC CAGCA CCCCAGCGAGCTCAGCAGTGACTGTGACAGCAGCG 3420
Db
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Db
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Qy
3541 GTTCTAGGAGCGAGGGCTATATTGTGGGGGGAATAAATCAGTTCAAGGGAAGTCTGGGAGA 3600
Db
3541 GTTCTAGGAGCGAGGGCTATATTGTGGGGGGAATAAATCAGTTCAAGGGAAGTCTGGGAGA 3600
Qy
3601 CTTGATTCTTAATACATATATTTTCTTTTCAAGCTGAGTAAATTTCTGAGCAAGTCAACAAG 3660
Db
3601 CTTGATTCTTAATACATATATTTTCTTTTCAAGCTGAGTAAATTTCTGAGCAAGTCAACAAG 3660
Qy
3661 GTAGTAACTGAGGCTGTAAAGATTACTTAGTTTCTCTTATTTAGGAATCTTTTTCTCTGT 3720
Db
3661 GTAGTAACTGAGGCTGTAAAGATTACTTAGTTTCTCTTATTTAGGAATCTTTTTCTCTGT 3720
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3721 GGAAGTTAGCAGCAAGGGCAATCCCGTTTCTTTTAAACAGGAAGAAACATTTCTTAAGAG 3780
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3721 GGAAGTTAGCAGCAAGGGCAATCCCGTTTCTTTTAAACAGGAAGAAACATTTCTTAAGAG 3780
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3781 TAAAGCCAAACAGATTCAAGCTTAGGCTTGTGCTGACTATATGATTGGTTTTTTCGAAAAAT 3840
Db
3781 TAAAGCCAAACAGATTCAAGCTTAGGCTTGTGCTGACTATATGATTGGTTTTTTCGAAAAAT 3840
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3841 CATTTCAGCGATGTTTACTATCTGATTCTAGAAATGAGACTAGTACCCTTTTGGTCAAGCTG 3900
Db
3841 CATTTCAGCGATGTTTACTATCTGATTCTAGAAATGAGACTAGTACCCTTTTGGTCAAGCTG 3900
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3901 TAAACAAACACCCAGTTGTAATGTCTCAAGTTTCAAGCTTAACTGCGAAGCAATCAAAA 3960
Db
3901 TAAACAAACACCCAGTTGTAATGTCTCAAGTTTCAAGCTTAACTGCGAAGCAATCAAAA 3960
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3961 AGAATAGAAATCTTTAGAGCAAACTGTGTTTTCTCCACATCTGGAGGTGAGTCTGCCAGGC 4020
Db
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4021 AGTTTGGAAATATTTACTTTCACAAAGTATTGACACTGTTTGTGTTTAAACAACTAAAGT 4080
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4021 AGTTTGGAAATATTTACTTTCACAAAGTATTGACACTGTTTGTGTTTAAACAACTAAAGT 4080
Qy
4081 TGCTCAAGGCAATCAATATTTCAAGTGGCTTAAAGTACTTCTGACAGTTTGGTATAT 4140
Db
4081 TGCTCAAGGCAATCAATATTTCAAGTGGCTTAAAGTACTTCTGACAGTTTGGTATAT 4140
Qy
4141 TTATTTGGCTATTGGCCATTTGTT 4200
Db
4141 TTATTTGGCTATTGGCCATTTGTT 4200
Qy
4201 GGAATTTAACTTACCTTACAGTCCAGAAAGCTGTGAAATTTGAATGAGGAAAAAATTAATTTT 4260
Db
4201 GGAATTTAACTTAACTTACAGTCCAGAAAGCTGTGAAATTTGAATGAGGAAAAAATTAATTTT 4260
Qy
4261 TGTTTTTACCACTTCTTAACTTAAATTTTAACTTAACTTAACTTAACTTAACTTAACTTAA 4320
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4321 CTCAAAGTGGTAAATACAGTCTGATTTTGTGCTTAACTTAACTTAACTTAACTTAACTTAA 4380
Db
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Qy
4381 TTATACTATATTACAGTCTTGTGAGTACGTTTGTAAAGTAAATTTTATATCTCAAACTA 4440
Db
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Qy
4441 CTTTGAATTTAGACCTCTGCTGGATCTTGTGTTTTTAACTTAACTTAACTTAACTTAACTTAA 4500
Db
4441 CTTTGAATTTAGACCTCTGCTGGATCTTGTGTTTTTAACTTAACTTAACTTAACTTAACTTAA 4500
Qy
4501 ATTTTGTATTTTGTATTAATCATATTTCATTATCTTCTTCTTCTTCTTCTTCTTCTTCTT 4560
Db
4501 ATTTTGTATTTTGTATTAATCATATTTCATTATCTTCTTCTTCTTCTTCTTCTTCTTCTT 4560
Qy
4561 ATATATTTGAAACATCTTTCTGAGAGAGTCTCCCGAGATTTCCCAATGAGGTTCTTGG 4620
Db
4561 ATATATTTGAAACATCTTTCTGAGAGAGTCTCCCGAGATTTCCCAATGAGGTTCTTGG 4620

	Query Match	100.0%	Score 5271;	DB 5;	Length 6169;
	Best Local Similarity	100.0%;	Pred. No. 0;	Mismatches	0;
	Matches 5271;	Conservative	0;		
QY	1	ATCTTTGTTCAGT	TACCTCAGGCGCTATTATGAATGAAATGAAATGAGATAACCAATGTGTAAG	60	
Db	1	ATCTTTGTTCAGT	TACCTCAGGCGCTATTATGAATGAAATGAAATGAGATAACCAATGTGTAAG	60	
QY	61	TCCTATAAATCTGTATAGCCTCCATTCGGATGTATGCTTTGGCAGGATGATAAGAATCA	120		
Db	61	TCCTATAAATCTGTATAGCCTCCATTCGGATGTATGCTTTGGCAGGATGATAAGAATCA	120		
QY	121	GGAAGAGAGATTCACAGTTTAGCCAAGTGTCCAGGCTGTCTGTCTCTTTATTATTAGTGA	180		
Db	121	GGAAGAGAGATTCACAGTTTAGCCAAGTGTCCAGGCTGTCTGTCTCTTTATTATTAGTGA	180		
QY	181	CAGATGTTGCTCCTGCAGACAGAGCTATTCTTCAGGAAACATCACATCAATATGTTAAATC	240		
Db	181	CAGATGTTGCTCCTGCAGACAGAGCTATTCTTCAGGAAACATCACATCAATATGTTAAATC	240		
QY	241	CATCAAAA CAGGAGCTTAAGAAAA CAGGAAATGAGATGGGCACTTGCCCAAAGSAAAAATGCCAG	300		
Db	241	CATCAAAA CAGGAGCTTAAGAAAA CAGGAAATGAGATGGGCACTTGCCCAAAGSAAAAATGCCAG	300		
QY	301	GAGAGCAAAATATGATGAAAAATAAACCTTTTCCCCTTTGTTTTTAAATTTTCAGHAAAAAATG	360		
Db	301	GAGAGCAAAATATGATGAAAAATAAACCTTTTCCCCTTTGTTTTTAAATTTTCAGHAAAAAATG	360		
QY	361	ATGAGGACCAAAATCAATGAAATAGGAAAAACAGCTCAGAAAAAAGATGTTTTCCAAATTTG	420		
Db	361	ATGAGGACCAAAATCAATGAAATAGGAAAAACAGCTCAGAAAAAAGATGTTTTCCAAATTTG	420		
QY	421	TAAATTAAGTATTGTTTCCCTTTGGGAGAGACCTCCCATGTGAGCTTTGATGGGAAAAATGGGAA	480		
Db	421	TAAATTAAGTATTGTTTCCCTTTGGGAGAGACCTCCCATGTGAGCTTTGATGGGAAAAATGGGAA	480		
QY	481	AAACGTCAAAAACATGATCTGATCAGATCCCAAGTGGATTTATTTTAAACACAGAT	540		
Db	481	AAACGTCAAAAACATGATCTGATCAGATCCCAAGTGGATTTATTTTAAACACAGAT	540		
QY	541	GSCATCACTCTGGGAGGAGCAAGTTTCAGGAGGTCTATGTTAGCAAGAGACATAACAATAAC	600		
Db	541	GSCATCACTCTGGGAGGAGCAAGTTTCAGGAGGTCTATGTTAGCAAGAGACATAACAATAAC	600		
QY	601	AGCAAAATCAAAAATTCGCAAAATGCAGGAGSAAAAATGGGACTGGGAAAAAGCTTTTATAAC	660		
Db	601	AGCAAAATCAAAAATTCGCAAAATGCAGGAGSAAAAATGGGACTGGGAAAAAGCTTTTATAAC	660		
QY	661	AGTGATTAGGAGTTCACATGTTTCGCAACACCTCCCCGTCTATACAGGAGACACAAAA	720		
Db	661	AGTGATTAGGAGTTCACATGTTTCGCAACACCTCCCCGTCTATACAGGAGACACAAAA	720		
QY	721	ATTGACTGGGCTTAGGCTTGAGCTTTCAAGGAGATATGAAAACTGAGAGCAAAACAAAA	780		
Db	721	ATTGACTGGGCTTAGGCTTGAGCTTTCAAGGAGATATGAAAACTGAGAGCAAAACAAAA	780		
QY	781	GACATGTTTAAAAAGGCAACACAGAACATTTGTGAGCCTTCAAGAGCAGCTGCCCCCAGCA	840		
Db	781	GACATGTTTAAAAAGGCAACACAGAACATTTGTGAGCCTTCAAGAGCAGCTGCCCCCAGCA	840		
QY	841	GGGACCCCTGAGGCATTTGCCCTTTAGGAAGGCCAGTTTTTCTTAAAGGATCTTTAAGAACTC	900		
Db	841	GGGACCCCTGAGGCATTTGCCCTTTAGGAAGGCCAGTTTTTCTTAAAGGATCTTTAAGAACTC	900		
QY	901	TTGAAAGATCATGAATTTTAAACCATTATTAAGTATAAAACAATATGCGATGATATATCAG	960		
Db	901	TTGAAAGATCATGAATTTTAAACCATTATTAAGTATAAAACAATATGCGATGATATATCAG	960		
QY	961	TTTAGACATGGGCTCCCAATTTTAAAGTACAGGATCAAGGATAACCGTGTCCCAGCTCC	1020		
Db	961	TTTAGACATGGGCTCCCAATTTTAAAGTACAGGATCAAGGATAACCGTGTCCCAGCTCC	1020		
QY	1021	GGATAGGTCAGAAATTCATTAGAAATCACTGTGTGGCATCTCTTAATCTTTTTCAGATGATC	1080		

US-10-244-633-3
; Sequence 3, Application US/10244633
; Publication No. US20030068640A1
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis,
; TITLE OF INVENTION: Prognosis And Treatment Of Glaucoma And Related
; FILE OF INVENTION: Disorders
; FILE OF INVENTION: 07425.0057.US01
; CURRENT APPLICATION NUMBER: US/10/244,633
; CURRENT FILING DATE: 2002-09-17
; PRIOR APPLICATION NUMBER: US/09/306,828
; PRIOR FILING DATE: 1999-05-07
; PRIOR APPLICATION NUMBER: US 09/227,881
; PRIOR FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 3
; LENGTH: 6169
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-244-633-3

Db 1021 |||||GGATAGGTCAGAAATCATTAGAAATCACTGTGTCCCATCTTAATCTTTTCAGAAATCAATC 1080
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Db 1081 TGTCTATAGCCCTCACAACAGGCGCGATGTGTCTGACCTACAACACATCTACAAACCAA 1140
Qy 1141 GTGCTCAACCAATTTAAACGTGTCTCATCTAGTAGTGTCCATTAACAATGCCACCTCCCC 1200
Db 1141 GTGCTCAACCAATTTAAACGTGTCTCATCTAGTAGTGTCCATTAACAATGCCACCTCCCC 1200
Qy 1201 TGTGAGCCCAATCCCGCTCCACAGGAAAGTCTCCCACTCTAGACTTCTGCAATCAGATGT 1260
Db 1201 TGTGAGCCCAATCCCGCTCCACAGGAAAGTCTCCCACTCTAGACTTCTGCAATCAGATGT 1260
Qy 1261 TACAGCCAGAAGCTCCGTAGGGTGTGCTGTCTTACCTACTACCTGTATGCTCTAC 1320
Db 1261 TACAGCCAGAAGCTCCGTAGGGTGTGCTGTCTTACCTACTACCTGTATGCTCTAC 1320
Qy 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGTCTCAGCCTCC 1380
Db 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGTCTCAGCCTCC 1380
Qy 1381 CGGTAGCTGGGACTACAGGCGCACGCGCGCTAAATTTTGTATTTGTAGTAGAGTGG 1440
Db 1381 CGGTAGCTGGGACTACAGGCGCACGCGCGCTAAATTTTGTATTTGTAGTAGAGTGG 1440
Qy 1441 GTTTACCAATATTAGCCGGTGTCTTGAACCTCTGACCTCAGGTGATCCACCACTC 1500
Db 1441 GTTTACCAATATTAGCCGGTGTCTTGAACCTCTGACCTCAGGTGATCCACCACTC 1500
Qy 1501 AGCTCTCAAGTCTGGGATACAGGCAATGAGTCAAGGCGCGCGCTCAAGGTCAGT 1560
Db 1501 AGCTCTCAAGTCTGGGATACAGGCAATGAGTCAAGGCGCGCGCTCAAGGTCAGT 1560
Qy 1561 TTAATAAGGAATAAATGAATGTTTAACTAAACCAACAGGAAACAGACAAAGCTGTGA 1620
Db 1561 TTAATAAGGAATAAATGAATGTTTAACTAAACCAACAGGAAACAGACAAAGCTGTGA 1620
Qy 1621 TAATTTCAAGGATTTCTGGGATGGGAAATGGTGCATGAGTGCCTGCTAGTCCAGAC 1680
Db 1621 TAATTTCAAGGATTTCTGGGATGGGAAATGGTGCATGAGTGCCTGCTAGTCCAGAC 1680
Qy 1681 CACTGTGCTCATCATCTTCTTCCCTCATCTCATTTTCAAGCTAAGTACCAATTTATT 1740
Db 1681 CACTGTGCTCATCATCTTCTTCCCTCATCTCATTTTCAAGCTAAGTACCAATTTATT 1740
Qy 1741 CACCATGCTTTTGTGTAAGCTCCACATCGTTACTGAAATTAAGAGTATACATAAATAG 1800
Db 1741 CACCATGCTTTTGTGTAAGCTCCACATCGTTACTGAAATTAAGAGTATACATAAATAG 1800
Qy 1801 TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGGAGGCAATACCCAGAGACTCCT 1860
Db 1801 TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGGAGGCAATACCCAGAGACTCCT 1860
Qy 1861 TGAAGCCCCGGCAGAGGTTTCTCTCCAGTGGGAGCCCTGCAAGACCCGGGTCC 1920
Db 1861 TGAAGCCCCGGCAGAGGTTTCTCTCCAGTGGGAGCCCTGCAAGACCCGGGTCC 1920
Qy 1921 TGGGTGCTCTGAGCAACCTGCAAGCCGTGCACTGTGTTTGTATCACTCTCTAGG 1980
Db 1921 TGGGTGCTCTGAGCAACCTGCAAGCCGTGCACTGTGTTTGTATCACTCTCTAGG 1980
Qy 1981 GACTGTGTCTTCTATTCTGTGTGACTGCTGTTTCAATTCATCCAGGCAATTCATTGACAAAT 2040
Db 1981 GACTGTGTCTTCTATTCTGTGTGACTGCTGTTTCAATTCATCCAGGCAATTCATTGACAAAT 2040
Qy 2041 TATTGAGTACTTATCTGCCAGACACAGGCAATGCTGAGCAAGCACTGCTC 2100
Db 2041 TATTGAGTACTTATCTGCCAGACACAGGCAATGCTGAGCAAGCACTGCTC 2100
Qy 2101 CCTACCTTCGTGGAGTGCAGATTTCTCATGGAAGCTGCAGAGAAATTAATAGCCA 2160
Db 2101 CCTACCTTCGTGGAGTGCAGATTTCTCATGGAAGCTGCAGAGAAATTAATAGCCA 2160

Db 2101 CCTACCTTCGTGGAGTGCAGATTTCTCATGGAAGCTGCAGAGAAATTAATAGCCA 2160
Qy 2161 GCCAACTTAACCCAGTGTGAAAGAAAGAAATAACACATCTTCAAGAAATTTGTGGC 2220
Db 2161 GCCAACTTAACCCAGTGTGAAAGAAAGAAATAACACATCTTCAAGAAATTTGTGGC 2220
Qy 2221 AGCATCCCTTAAACAAGGCCACCTCCCTAGCGCCCTGCTGCTCTCCTCATCTGTCGCCGAGG 2280
Db 2221 AGCATCCCTTAAACAAGGCCACCTCCCTAGCGCCCTGCTGCTCTCCTCATCTGTCGCCGAGG 2280
Qy 2281 CCCCCAAGCCGAGTCTTCAAGCCTCTCCTCATCAGTCAACAGCCTCAGCTGAGCT 2340
Db 2281 CCCCCAAGCCGAGTCTTCAAGCCTCTCCTCATCAGTCAACAGCCTCAGCTGAGCT 2340
Qy 2341 GCCTCGTCTCCCTGTGATCTGTGTGATCTGTGTGATCTGTGTGATCTGTGTGATCTGTGTG 2400
Db 2341 GCCTCGTCTCCCTGTGATCTGTGTGATCTGTGTGATCTGTGTGATCTGTGTGATCTGTGTG 2400
Qy 2401 CCAGAAAGGAAATGGAGAGGAAATAGTCTAAACGGAGAAATCTGGAGGGACAGTGTTC 2460
Db 2401 CCAGAAAGGAAATGGAGAGGAAATAGTCTAAACGGAGAAATCTGGAGGGACAGTGTTC 2460
Qy 2461 CTCAAGAGGAAAGGGGCTCCAGTCCAGGAGAAATTCAGAGAGTGGGACCTGAGGAG 2520
Db 2461 CTCAAGAGGAAAGGGGCTCCAGTCCAGGAGAAATTCAGAGAGTGGGACCTGAGGAG 2520
Qy 2521 TGGGAGCTTGGGCGCTGAGCGGCTGCTGAAAGCAGGAGGTAAGAGGCGCAAGGCTGAA 2580
Db 2521 TGGGAGCTTGGGCGCTGAGCGGCTGCTGAAAGCAGGAGGTAAGAGGCGCAAGGCTGAA 2580
Qy 2581 GCTGCCAGATGTTCAAGTGTGTTTCAAGCGGCTGGAGTGTTCCTGTTCTCTGTGAGC 2640
Db 2581 GCTGCCAGATGTTCAAGTGTGTTTCAAGCGGCTGGAGTGTTCCTGTTCTCTGTGAGC 2640
Qy 2641 CTTTTTATCTTTCTCTGCTTGGAGAGAAAGTCTATTTCATGAAGGATGAGTGTTC 2700
Db 2641 CTTTTTATCTTTCTCTGCTTGGAGAGAAAGTCTATTTCATGAAGGATGAGTGTTC 2700
Qy 2701 ATAAAGTCACTGTTAAATTCAGGCTGTCAGTGTGTTTCTTCTTCAAGAGGCTTTAT 2760
Db 2701 ATAAAGTCACTGTTAAATTCAGGCTGTCAGTGTGTTTCTTCTTCAAGAGGCTTTAT 2760
Qy 2761 TTAATGGGAATATAGGAAGCGAGCTCAATTTCTTAGGCGGTAAATTCACGGAAGAGTGAC 2820
Db 2761 TTAATGGGAATATAGGAAGCGAGCTCAATTTCTTAGGCGGTAAATTCACGGAAGAGTGAC 2820
Qy 2821 TGGAGTCTTTTCTTTTCTGCTTGGGCAACTACTCAGCCCTGTGTGTGAGTGTGCTTA 2880
Db 2821 TGGAGTCTTTTCTTTTCTGCTTGGGCAACTACTCAGCCCTGTGTGTGAGTGTGCTTA 2880
Qy 2881 TGCAGACCGTTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTCTGTTCTGCCATT 2940
Db 2881 TGCAGACCGTTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTCTGTTCTGCCATT 2940
Qy 2941 GGTGCTGTGCGACCGTGGGCAAGTGTCTCTCTTCCCTGGGCAACTACTCAGCTCTGCT 3000
Db 2941 GGTGCTGTGCGACCGTGGGCAAGTGTCTCTCTTCCCTGGGCAACTACTCAGCTCTGCT 3000
Qy 3001 ATAAAGACCTTGCAGCTCTCGTGTGTCTGTGAACACTTCCCTGTGATTTCTGTGAGGG 3060
Db 3001 ATAAAGACCTTGCAGCTCTCGTGTGTCTGTGAACACTTCCCTGTGATTTCTGTGAGGG 3060
Qy 3061 GATGTTGAGAGGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3120
Db 3061 GATGTTGAGAGGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3120
Qy 3121 GAGACGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3180
Db 3121 GAGACGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3180
Qy 3181 CAGGACCGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3240
Db 3181 CAGGACCGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3240

Qy	3241	TCCCTAAGCATAGACAATGGCAATTTGGCCAATAACCAAAAAAGAAATGCAGAGACTAACTGGT	3300
Db	3241		
Qy	3301	GGTAGCTTTTGGCTTGGCAATTCAAAAA CTGGGCGAGAGCAAGTGAGAAATGCCAGAGATTG	3360
Db	3301		
Qy	3361	TTAAACTTTTTCACCCTGACGAGCACCCACGCGAGCTCAGCAGTGTACTGCTGACAGCACGG	3420
Db	3361		
Qy	3421	AGTGACCTTCGACGCGCAGGGGAGGAGAAAGAGAGAGGGATAGTGTATGAGCAAGAAAG	3480
Db	3421		
Qy	3481	ACAGATTCAATTCAGGGGAGTGGGAATTGACCA CAGGGATATAGTCCACGTGATCCTGG	3540
Db	3481		
Qy	3541	GTTCTAGGAGCAGGGCTATATTGTGGGGGAAAAAATCAGTTCAAGGGAAGTGCGGAGA	3600
Db	3541		
Qy	3601	CCTGATTTCTAATACTATA TTTTTCCTTTTCAAGCTGAGTAA TCTTGAGCAAGTCAACAG	3660
Db	3601		
Qy	3661	GTAGTAAGTGGGCTGTGAAGATTACTAGTTTCTCCCTATTAGGAACTCTTTTCTCTGT	3720
Db	3661		
Qy	3721	GGAGTTAGCAGCA CAAAGGGCAATCCCGTTTCTTTTAA CAGGAAGAAACATTCCTAAGAG	3780
Db	3721		
Qy	3781	TAAAGCCAAACAGATTCAGGCTTAGGCTTGTCTGACTATATGATTTGGTTTTTTGAAAAAT	3840
Db	3781		
Qy	3841	CATTTTCAGCGATGTTTACTATCTCATTT CAGAAAATGAGACTAGTACCCCTTTGGTCAGCTG	3900
Db	3841		
Qy	3901	TAAACAAACACCCAGTTGTAAATGCTCAAGTTCAGGCTTAACTGCAGAACCAATCAAAA	3960
Db	3901		
Qy	3961	AGAAATAGAAATCTTTTAGAGCAAACTGTGTTTCTCCACATCTGGAGGTGAGTCTGCCAGGGC	4020
Db	3961		
Qy	4021	AGTTTGGAAATATTTACTTTCACAAGTATTGACACTGTTTGGTATTAA CAA CATAAAGT	4080
Db	4021		
Qy	4081	TGCTCAAGGCAATCATTTATTTCAAGTGGCTTAAAGTTACTTCTGACAGTTTTGGTATAT	4140
Db	4081		
Qy	4141	TTATTGGCTATTGCCATTTGCTTTTTTTGTTTTTCTCTTGGGTTTATTAATGTAAAGCAG	4200
Db	4141		
Qy	4201	GGATTATTAACTACAGTCCAGAAAGCCTGTGAATTTTGAATTGAGAAAAAATTACATTTT	4260
Db	4201		
Qy	4261	TGTTTTTACCACTTCTAACTAAATTTTAA CATTTTTCCATTTCGGAATAGAGCCATAAA	4320
Db	4261		

Qy	4321	CTCAAAGTGGTAATAACACGTAACCTGTGTGATTTTGTCATTACCAAATAGANAATCAAGACATT	4380
Db	4321	CTCAAAGTGGTAATAAACACGTAACCTGTGTGATTTTGTCATTACCAAATAGANAATCAAGACATT	4380
Qy	4381	TTATACTATATTAACAGTTGTGCAGATACGTTCTAAGTGAADATATTTATATCTCAAACCTA	4440
Db	4381	TTATACTATATTAACAGTTGTGCAGATACGTTCTAAGTGAADATATTTATATCTCAAACCTA	4440
Qy	4441	CTTTGAAAATPAGACCTCCTGCTGGATCTTGTTTTTTAAACATATTAATAACAATGTTTAA	4500
Db	4441	CTTTGAAAATPAGACCTCCTGCTGGATCTTGTTTTTTAAACATATTAATAACAATGTTTAA	4500
Qy	4501	ATTTTGATATTTTGAATAATCATATTTTCATATCATTTGTTTCCCTTTGTGTAATCTATATTTT	4560
Db	4501	ATTTTGATATTTTGAATAATCATATTTTCATATCATTTGTTTCCCTTTGTGTAATCTATATTTT	4560
Qy	4561	ATATNTTGAARAACATCTTTCTCAGAAGAGTTCCCCAGATTTTCCACCAATGAGGTTCTTGG	4620
Db	4561	ATATNTTGAARAACATCTTTCTCAGAAGAGTTCCCCAGATTTTCCACCAATGAGGTTCTTGG	4620
Qy	4621	CATGCAACACACAGAGTAGAACTGATTTTAGAGGCTAAACATTTGACATTTGGTGCCTGAGA	4680
Db	4621	CATGCAACACACAGAGTAGAACTGATTTTAGAGGCTAAACATTTGACATTTGGTGCCTGAGA	4680
Qy	4681	TGCAAGACTGAAATTTAGAAAGTTCTCCCAAAGATACACAGTTGTTTAAAGCTAGGGGTG	4740
Db	4681	TGCAAGACTGAAATTTAGAAAGTTCTCCCAAAGATACACAGTTGTTTAAAGCTAGGGGTG	4740
Qy	4741	AGGGGGGAATCTCGCGCTTCATAGGAATGCTCTCCCTGGAGCCTGGTAGGGTCTGTCTC	4800
Db	4741	AGGGGGGAATCTCGCGCTTCATAGGAATGCTCTCCCTGGAGCCTGGTAGGGTCTGTCTC	4800
Qy	4801	CTTGTGTTCTGGCTGGCTGTATTTTTTCTCTGCTCCCTGCTACGTCCTTTAAAGGACTTGT	4860
Db	4801	CTTGTGTTCTGGCTGGCTGTATTTTTTCTCTGCTCCCTGCTACGTCCTTTAAAGGACTTGT	4860
Qy	4861	GGATCTCCAGTTCCTAGCATAGTGCCTGGCACAGTGCAGGTTCTCAATGAGTTTGACAG	4920
Db	4861	GGATCTCCAGTTCCTAGCATAGTGCCTGGCACAGTGCAGGTTCTCAATGAGTTTGACAG	4920
Qy	4921	TGAATGGAATATAAACTAGAAATATATCTCTGTGTAATTCAGACACACCAAGTAGTCTCG	4980
Db	4921	TGAATGGAATATAAACTAGAAATATATCTCTGTGTAATTCAGACACACCAAGTAGTCTCG	4980
Qy	5041	TAGAACTATTAATGGGGTATGGGTGCATAAATTTGGGATGTTCTTTTAAAAAGAACTC	5100
Db	5041	TAGAACTATTAATGGGGTATGGGTGCATAAATTTGGGATGTTCTTTTAAAAAGAACTC	5100
Qy	5101	CAAAACAGCTTCGGAAGGTATTTTCTAAGATCTTGTGGCAGCGTGAAGCAACCCC	5160
Db	5101	CAAAACAGCTTCGGAAGGTATTTTCTAAGATCTTGTGGCAGCGTGAAGCAACCCC	5160
Qy	5161	CCTGTGCACAGCCCCACCCAGCTCAGTGGCCACCTCTCTTCCCCCATGAAGGCGTG	5220
Db	5161	CCTGTGCACAGCCCCACCCAGCTCAGTGGCCACCTCTCTTCCCCCATGAAGGCGTG	5220
Qy	5221	GCTCCCCAGTATATATAAACCTCTCTGGAGCTCGGGCATGAGCCAAGG	5271
Db	5221	GCTCCCCAGTATATATAAACCTCTCTGGAGCTCGGGCATGAGCCAAGG	5271

RESULT 3
US-09-985-637A-1
; Sequence 1, Application US/09985637A
; Publication No. US2003011900A1
; GENERAL INFORMATION:
; APPLICANT: Polansky, Jon
; TITLE OF INVENTION: METHODS TO SCREEN
; TITLE OF INVENTION: METHODS TO DEVELOP GLA
; TITLE OF INVENTION: TO DEVELOP GLA

FILE REFERENCE: 13587.296
CURRENT APPLICATION NUMBER: US/09/985,637A
CURRENT FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 21
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 5300
TYPE: DNA
ORGANISM: Homo sapiens
US-09-985-637A-1

Query Match 99.5%; Score 5246.4; DB 3; Length 5300;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 5269; Conservative 0; Mismatches 1; Indels 2; Gaps 2;

QY	1	ATCTTTGTTCCAGTTTACCTCAGGGCTATTATGAAATGAAATGAGATACCAATGTGAAG	60
DB	1	ATCTTTGTTCCAGTTTACCTCAGGGCTATTATGAAATGAAATGAGATACCAATGTGAAG	60
QY	61	TCCTATAAATCTGTATAGCCTCCATTCGGATGTATGCTTTGGCAGGATGATAAAGATCA	120
DB	61	TCCTATAAATCTGTATAGCCTCCATTCGGATGTATGCTTTGGCAGGATGATAAAGATCA	120
QY	121	GGAAAGAGAGTAGTCCACGTTAGCCAAAGTGTCCAGGCTGTGTCTCTTATTTAGTGA	180
DB	121	GGAAAGAGAGTAGTCCACGTTAGCCAAAGTGTCCAGGCTGTGTCTCTTATTTAGTGA	180
QY	181	CAGATGTGCTCTGTAGCAGAAAGCTATTTCTCAGAAACATCAATCCAAATATGGTAAATC	240
DB	181	CAGATGTGCTCTGTAGCAGAAAGCTATTTCTCAGAAACATCAATCCAAATATGGTAAATC	240
QY	241	CATCAACAGGAGCTAAGAAACAGGATGAGTGGCAGCTTCCAGGAAATATGCCAG	300
DB	241	CATCAACAGGAGCTAAGAAACAGGATGAGTGGCAGCTTCCAGGAAATATGCCAG	300
QY	301	GAGAGCAAAATTAATCATGATAAAATAAATTTTCCCTTTTGTGTTTAAATTTAGGAAATATG	360
DB	301	GAGAGCAAAATTAATCATGATAAAATAAATTTTCCCTTTTGTGTTTAAATTTAGGAAATATG	360
QY	361	ATGAGGACCAAAATCAATGATTAAGGAAACAGCTCAGAAAAAAGATGTTTCCAAATGG	420
DB	361	ATGAGGACCAAAATCAATGATTAAGGAAACAGCTCAGAAAAAAGATGTTTCCAAATGG	420
QY	421	TAATTAAGTATTTGTTCTTGGAGAGACCTCCATGTGAGCTTGATGGAAATCGAA	480
DB	421	TAATTAAGTATTTGTTCTTGGAGAGACCTCCATGTGAGCTTGATGGAAATCGAA	480
QY	481	AAACGTCAAAAGCATGATCTGATCAGATCCCAAAGTGGATTATTTTAAAAACCCAGAT	540
DB	481	AAACGTCAAAAGCATGATCTGATCAGATCCCAAAGTGGATTATTTTAAAAACCCAGAT	540
QY	541	GGCATCACTTGGGGAGGCAAGTTTCAGGAAGGTCATGTTAGCAAAAGCATATAAATAAC	600
DB	541	GGCATCACTTGGGGAGGCAAGTTTCAGGAAGGTCATGTTAGCAAAAGCATATAAATAAC	600
QY	601	AGCAAAATCAAAATTCGCAAAATGACAGGAGGAAATGGGACCTGGGAAAGCTTTTCAATAC	660
DB	601	AGCAAAATCAAAATTCGCAAAATGACAGGAGGAAATGGGACCTGGGAAAGCTTTTCAATAC	660
QY	661	AGTGATTAGGACGTTGACCATGTTTCGCAACACCTCCCGCTCATATACAGGAAACACAAA	720
DB	661	AGTGATTAGGACGTTGACCATGTTTCGCAACACCTCCCGCTCATATACAGGAAACACAAA	720
QY	721	ATTGACTGGGCTAAGCTGTGACTTTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA	780
DB	721	ATTGACTGGGCTAAGCTGTGACTTTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA	780
QY	781	GACATGTTTAAAGGCAACAGAAATTTGAGCCTTCAAGCAGAGTGCCTCAGCA	840
DB	781	GACATGTTTAAAGGCAACAGAAATTTGAGCCTTCAAGCAGAGTGCCTCAGCA	840
QY	841	GGGACCCCTGAGGATTTGCTTTAGGAAGCCAGTGTCTTAAAGGAATCTTTAAGAACTC	900
DB			

DB	841	GGGACCCCTGAGGCAATTTGCTTTAGGAAGCCAGTGTCTTAAAGGAATCTTTAAGAACTC	900
QY	901	TTGAAAGATCATGAATTTTAAACATTTTAAGTATAAAACAATATATGCGATGCAATACAG	960
DB	901	TTGAAAGATCATGAATTTTAAACATTTTAAAGTATAAAACAATATATGCGATGCAATACAG	960
QY	961	TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAAGGATAACGTGTCCAGCTCC	1020
DB	961	TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAAGGATAACGTGTCCAGCTCC	1020
QY	1021	GGATAGGTGAGAAATCATTTAGAAATCACTGTGTGCCCATCTTAACCTTTTTCAGAAATGATC	1080
DB	1021	GGATAGGTGAGAAATCATTTAGAAATCACTGTGTGCCCATCTTAACCTTTTTCAGAAATGATC	1080
QY	1081	TGTCATAGCCCTCACACACAGGCCCGATGTGTCTGACCTTACACACATCTACAACCAA	1140
DB	1081	TGTCATAGCCCTCACACACAGGCCCGATGTGTCTGACCTTACACACATCTACAACCAA	1140
QY	1141	GTGCTTCAACCATTTGTAAAGTGTCTCATCTCAGTAGGTCCCATTTACAAATGCCACCTCC	1200
DB	1141	GTGCTTCAACCATTTGTAAAGTGTCTCATCTCAGTAGGTCCCATTTACAAATGCCACCTCC	1200
QY	1201	TGTGACGCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTTCTGCATCACGATGT	1260
DB	1201	TGTGACGCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTTCTGCATCACGATGT	1260
QY	1261	TACAGCCAGAGTCCGTGAGGCTGTGTCTTACACCTACCTGTGTGTCTCTAC	1320
DB	1261	TACAGCCAGAGTCCGTGAGGCTGTGTCTTACACCTACCTGTGTGTCTCTAC	1320
QY	1321	ACCTGAGCTCACTGCAACCTCTGCCCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC	1380
DB	1321	ACCTGAGCTCACTGCAACCTCTGCCCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC	1380
QY	1381	CGCGTAGCTGGGACTACAGGCGCACGCCGGCTAAATTTTGTATTTAGTAGAGATGGG	1440
DB	1381	CGCGTAGCTGGGACTACAGGCGCACGCCGGCTAAATTTTGTATTTAGTAGAGATGGG	1440
QY	1441	GTTCACCATATTTAGCCGGCTGTGTGAACTCTGACCTCAGGTGATCCACCCCTC	1500
DB	1441	GTTCACCATATTTAGCCGGCTGTGTGAACTCTGACCTCAGGTGATCCACCCCTC	1500
QY	1501	AGCCTCTTAAAGTGTGGGATTTACAGGATGAGTCAACCGCCCGCCAGGGTCAAGTGT	1560
DB	1501	AGCCTCTTAAAGTGTGGGATTTACAGGATGAGTCAACCGCCCGCCAGGGTCAAGTGT	1560
QY	1561	TTAATAAGGAATTAACCTTGAATGTTTACTTAAACCAACAGGAAACACAGAAAGCTGTGA	1620
DB	1561	TTAATAAGGAATTAACCTTGAATGTTTACTTAAACCAACAGGAAACACAGAAAGCTGTGA	1620
QY	1621	TAATTTTACGGGATTTCTGGGATGGGAAATGGTCCATGAGCTGCTGCTAGTCCAGAC	1680
DB	1621	TAATTTTACGGGATTTCTGGGATGGGAAATGGTCCATGAGCTGCTGCTAGTCCAGAC	1680
QY	1681	CACCTGCTCATCACTTTCTTCCCTCATCTCTTCTTTCAGGCTAAGTTTACCATTTTAT	1740
DB	1681	CACCTGCTCATCACTTTCTTCCCTCATCTCTTCTTTCAGGCTAAGTTTACCATTTTAT	1740
QY	1741	CACCATGCTTTTGTGGTAAAGCTCCACATCGTTACTGAAATAAGAGTATACATAAATAG	1800
DB	1741	CACCATGCTTTTGTGGTAAAGCTCCACATCGTTACTGAAATAAGAGTATACATAAATAG	1800
QY	1801	TTTCATTTTGGGCTCATCTGTGTGTGTATAGGGAGGAGGCGATACCCAGAGACTCT	1860
DB	1801	TTTCATTTTGGGCTCATCTGTGTGTGTATAGGGAGGAGGCGATACCCAGAGACTCT	1860
QY	1861	TGAAGCCCGCCGACAGAGTTCCTCTCCAGCTGGGGAGCCCTGCAAGCACCCGGGCTCC	1920
DB	1861	TGAAGCCCGCCGACAGAGTTCCTCTCCAGCTGGGGAGCCCTGCAAGCACCCGGGCTCC	1920
QY	1921	TGGGTGTCTGAGCAACCTGCCAGCCCGCTGCGACTGGTGTGTTTGTATCTCTCTAGG	1980
DB	1921	TGGGTGTCTGAGCAACCTGCCAGCCCGCTGCGACTGGTGTGTTTGTATCTCTCTAGG	1980

QY	1981	GACCTGTTGCTTTCTATTTCTGTGTGACTGTTCTATTCATTCATCCAGGCAATTCATGACAAAT	2040
DB	1981	GACCTGTTGCTTTCTATTTCTGTGTGACTGTTCTATTCATTCATCCAGGCAATTCATGACAAAT	2040
QY	2041	TATTGAGTACTTATATCTGCCACACACACAGAGACAAATGGTGTAGCAAGCAGTCACTGC	2100
DB	2041	TATTGAGTACTTATATCTGCCACACACACAGAGACAAATGGTGTAGCAAGCAGTCACTGC	2100
QY	2101	CCTACCTCTGTGAGGTGACAGTTTCTCATGGAAGACGTGCAGAGAAATTAATAGCCA	2160
DB	2101	CCTACCTCTGTGAGGTGACAGTTTCTCATGGAAGACGTGCAGAGAAATTAATAGCCA	2160
QY	2161	GCACACTTAAACCCAGTGTGAAGAAAGAAATAAACACCATCTTGAAGAAATTTGGCG	2220
DB	2161	GCACACTTAAACCCAGTGTGAAGAAAGAAATAAACACCATCTTGAAGAAATTTGGCG	2220
QY	2221	AGCATCCCTTAAACAGGCCACTCCCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGG	2280
DB	2221	AGCATCCCTTAAACAGGCCACTCCCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGG	2280
QY	2281	CCCCAAGCCCGAGTCTTCCAGCTCTCTCCATCAGTCAAGCGCTGCAGCTGGCCT	2340
DB	2281	CCCCAAGCCCGAGTCTTCCAGCTCTCTCCATCAGTCAAGCGCTGCAGCTGGCCT	2340
QY	2341	GCCTCGCTTCCCGTGAATCGTCTGTGTGACTCTGAGCTGGAGACTCTTGGCTCCAGGCT	2400
DB	2341	GCCTCGCTTCCCGTGAATCGTCTGTGTGACTCTGAGCTGGAGACTCTTGGCTCCAGGCT	2400
QY	2401	CCAGAAAGAAATCGAGAGGAACTAGTCTTAACGGAGATCTGAGGGGACAGTCTTTC	2460
DB	2401	CCAGAAAGAAATCGAGAGGAACTAGTCTTAACGGAGATCTGAGGGGACAGTCTTTC	2460
QY	2461	CTCAGAGGAAAGGGGCTCCAGTCCAGAGAAATCCAGGAGGTGGGACTCAGGGAG	2520
DB	2461	CTCAGAGGAAAGGGGCTCCAGTCCAGAGAAATCCAGGAGGTGGGACTCAGGGAG	2520
QY	2521	TGGGACGCTGGGCTGAGCGGTGCTGAAAGGAGAGAGGTGAAGGGCAAGGCTGAA	2580
DB	2521	TGGGACGCTGGGCTGAGCGGTGCTGAAAGGAGAGAGGTGAAGGGCAAGGCTGAA	2580
QY	2581	GCTGCCAGATGTTCAAGTGTGTTTACGGGCTGGAGTTTCCGTTCTCTGTGAGC	2640
DB	2581	GCTGCCAGATGTTCAAGTGTGTTTACGGGCTGGAGTTTCCGTTCTCTGTGAGC	2640
QY	2641	CTTTTATCTTTTCTCTGTTGAGGAGAGAGTCTATTTTCATGAGGAGTCAAGTTTC	2700
DB	2641	CTTTTATCTTTTCTCTGTTGAGGAGAGAGTCTATTTTCATGAGGAGTCAAGTTTC	2700
QY	2701	ATAAGTCAGCTGTTTAAATTCAGGGTGTGCATFGGTTTCTTCCAGAGGCTTTAT	2760
DB	2701	ATAAGTCAGCTGTTTAAATTCAGGGTGTGCATFGGTTTCTTCCAGAGGCTTTAT	2760
QY	2761	TAAATGGGATATAGGAGCGAGTCTATTTCTAGGCGGTTAATTCACGGAAGGTGAC	2820
DB	2761	TAAATGGGATATAGGAGCGAGTCTATTTCTAGGCGGTTAATTCACGGAAGGTGAC	2820
QY	2821	TGAGTCTTTTCTTTCTGTTCTGCGCACTACTCAGCGCTGTGTGGACTTTGGCTTA	2880
DB	2821	TGAGTCTTTTCTTTCTGTTCTGCGCACTACTCAGCGCTGTGTGGACTTTGGCTTA	2880
QY	2881	TGCAAGACGTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTTCTGTTCTGCCATT	2940
DB	2881	TGCAAGACGTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTTCTGTTCTGCCATT	2940
QY	2941	GGTTGGCTGTGCNACCGTGGCAGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	3000
DB	2941	GGTTGGCTGTGCNACCGTGGCAGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	3000
QY	3001	ATAAAGACCCCTGTCAGCTCTGCTGTTCTGTGAAACACTTCCCTGTGATTTCTCTGTGAGGG	3060
DB	3001	ATAAAGACCCCTGTCAGCTCTGCTGTTCTGTGAAACACTTCCCTGTGATTTCTCTGTGAGGG	3060
QY	3061	GGATCTTCCAGGGGAGGAGGAGCTGGAGCAGCTGAGCCACAGGGAGGTGGAGGG	3120
DB	3061	GGATCTTCCAGGGGAGGAGGAGCTGGAGCAGCTGAGCCACAGGGAGGTGGAGGG	3120
QY	3121	GGACAGGAGGAGGAGGAGGAGCTGGGTGCTCCATCAGTCTCTCACTGATCACGTCAGACTC	3180
DB	3121	GGACAGGAGGAGGAGGAGGAGCTGGGTGCTCCATCAGTCTCTCACTGATCACGTCAGACTC	3180
QY	3181	CAGGACCGAGGAGGAGGAGGAGCTTCAAGTCAAGTCAAGTCAAGTCAAGTCAAGTCAAGT	3240
DB	3181	CAGGACCGAGGAGGAGGAGGAGCTTCAAGTCAAGTCAAGTCAAGTCAAGTCAAGTCAAGT	3240
QY	3241	TCCCTTAAGCATAGACATAGGCAATTTGCCAATTAACCAAAAGAAATGAGAGACTTAACGTGT	3300
DB	3241	TCCCTTAAGCATAGACATAGGCAATTTGCCAATTAACCAAAAGAAATGAGAGACTTAACGTGT	3300
QY	3301	GGTAGCTTTTGGCTGGCATTTCAAAAACCTGGGCGAGAGCAAGTGGAAAAATGCCAGAGATTG	3360
DB	3301	GGTAGCTTTTGGCTGGCATTTCAAAAACCTGGGCGAGAGCAAGTGGAAAAATGCCAGAGATTG	3360
QY	3361	TTAAACCTTTTCAACCTGACAGCACCCCGAGCTCAGCAGCTCAGCAGCTCAGCAGCTCAGC	3420
DB	3361	TTAAACCTTTTCAACCTGACAGCACCCCGAGCTCAGCAGCTCAGCAGCTCAGCAGCTCAGC	3420
QY	3421	AGTGAACCTTCAACCTGACAGCACCCCGAGCTCAGCAGCTCAGCAGCTCAGCAGCTCAGC	3480
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QY	3481	ACAGATTTCAATCAAGGCGAGTGGGAATTCACACAGGGAATTAAGTCCACGTGATCCTGG	3540
DB	3481	ACAGATTTCAATCAAGGCGAGTGGGAATTCACACAGGGAATTAAGTCCACGTGATCCTGG	3540
QY	3541	GTTCTAGAGGCGAGGCTATATTTGGGGGAAAAAATCAGTTCAAGGAGAGTCCGGAGA	3600
DB	3541	GTTCTAGAGGCGAGGCTATATTTGGGGGAAAAAATCAGTTCAAGGAGAGTCCGGAGA	3600
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QY	3661	GTAGTAACCTGAGGCTGTAAAGATTACTAGTTCTCTTATAGGAATCTTTTCTCTCTGT	3720
DB	3661	GTAGTAACCTGAGGCTGTAAAGATTACTAGTTCTCTTATAGGAATCTTTTCTCTCTGT	3720
QY	3721	GGATTTAGCAGCACAAAGGCAATCCGTTCTTTTAAACAGGAGAAATCATTCCTAAGAG	3780
DB	3721	GGATTTAGCAGCACAAAGGCAATCCGTTCTTTTAAACAGGAGAAATCATTCCTAAGAG	3780
QY	3781	TAAAGCCAAACAGATTTCAAGCCTTAGCTCTTCTGACTATATGATTGGTTTTTTGAAAAAT	3840
DB	3781	TAAAGCCAAACAGATTTCAAGCCTTAGCTCTTCTGACTATATGATTGGTTTTTTGAAAAAT	3840
QY	3841	CATTTTCAGCGATGTTTACTATCTGATTTCAAGAAATGAGACTAGTACCTTTGGTCAAGTG	3900
DB	3841	CATTTTCAGCGATGTTTACTATCTGATTTCAAGAAATGAGACTAGTACCTTTGGTCAAGTG	3900
QY	3901	TAAACAAACACCCAGTTGTAATGCTCAAGTTCAAGCTTAAGTCAAGCAACCAATCAAA	3959
DB	3901	TAAACAAACACCCAGTTGTAATGCTCAAGTTCAAGCTTAAGTCAAGCAACCAATCAAA	3960
QY	3960	TAAGATAGAAATCTTTAGAGCAAACTGTGTTTCTCCACATCTGGAGGTGAGTCTGCCAGGG	4019
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QY	4020	CAGTTTGGAAATATTTTACTTCAAGTATGACACTGTTGTTGGTATTAACAAATAAAG	4079
DB	4020	CAGTTTGGAAATATTTTACTTCAAGTATGACACTGTTGTTGGTATTAACAAATAAAG	4079
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RESULT 4
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; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis,
; TITLE OF INVENTION: Prognosis And Treatment Of Glaucoma And Related
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: 07425.0057.US01
; CURRENT APPLICATION NUMBER: US/10/244,633
; CURRENT FILING DATE: 2002-09-17
; PRIOR APPLICATION NUMBER: US/09/306,828
; PRIOR FILING DATE: 1999-05-07
; PRIOR APPLICATION NUMBER: US 09/227,881
; PRIOR FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 1
; LENGTH: 5300
; TYPE: DNA
; ORGANISM: Homo sapiens
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 Qy |||||
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 Db |||||
 1981 GACCTGTGCTTTCTATTTCTGTGTGACTCGTTTCACTCAGGCAATTCATGACATTT 2040
 Qy |||||
 2041 TATTGAGTACTTATCTGCCAGACACAGAGCAAAATGTGAGCAAGAGCAAGTCACTGC 2100
 Db |||||
 2041 TATTGAGTACTTATCTGCCAGACACAGAGCAAAATGTGAGCAAGAGCAAGTCACTGC 2100
 Qy |||||
 2101 CCTACCTTCGTGGAGGTGACAGTTTCTCATGGAAGACGTGAGAGGAAATTAATADCCA 2160
 Db |||||
 2101 CCTACCTTCGTGGAGGTGACAGTTTCTCATGGAAGACGTGAGAGGAAATTAATADCCA 2160
 Qy |||||
 2161 GCCAACTTAAACCCAGTCTGAAGAAAGGAAATAACACCATCTTGAAGAAATTTGTGCG 2220
 Db |||||
 2161 GCCAACTTAAACCCAGTCTGAAGAAAGGAAATAACACCATCTTGAAGAAATTTGTGCG 2220
 Qy |||||
 2221 AGCATCCCTTAAACAGGCGACCTCCCTAGCGCCCTGCTGCTCCATCGTGGCCGAGG 2280
 Db |||||
 2221 AGCATCCCTTAAACAGGCGACCTCCCTAGCGCCCTGCTGCTCCATCGTGGCCGAGG 2280
 Qy |||||
 2281 CCCCAGAGCCGAGTCTTCAAGGCTCCTCTCCATCAGTCAACAGCGCTGAGCTGGCCT 2340
 Db |||||
 2281 CCCCAGAGCCGAGTCTTCAAGGCTCCTCTCCATCAGTCAACAGCGCTGAGCTGGCCT 2340
 Qy |||||
 2341 GCCTCGCTCCCGTGAATCGTCTGGTGATCTGAGCTGAGAGTCTCTTGGCTCCAGCT 2400
 Db |||||
 2341 GCCTCGCTCCCGTGAATCGTCTGGTGATCTGAGCTGAGAGTCTCTTGGCTCCAGCT 2400
 Qy |||||
 2401 CCAGAAAGGAATCGAGAGGGAATAGTCTTAAACGAGAACTCTGAGGGGACAGTGTTC 2460
 Db |||||
 2401 CCAGAAAGGAATCGAGAGGGAATAGTCTTAAACGAGAACTCTGAGGGGACAGTGTTC 2460
 Qy |||||
 2461 CTCAGAGGAAAGGGGCTCCACGTCCAGGAGAAATTCAGGAGTGGGGAGCTGAGGGAG 2520
 Db |||||
 2461 CTCAGAGGAAAGGGGCTCCACGTCCAGGAGAAATTCAGGAGTGGGGAGCTGAGGGAG 2520
 Qy |||||
 2521 TGGGGACGCTGGGGCTGAGCGGGTGTCTGAAGGAGGAGAGGTGAAAGGGCAGGCTGAA 2580
 Db |||||
 2521 TGGGGACGCTGGGGCTGAGCGGGTGTCTGAAGGAGGAGAGGTGAAAGGGCAGGCTGAA 2580

2581 GCTCCAGATGTTTCAGTGTGTTTTCAGGGGCTGGAGTTTTTCGTTCTCTCTCTGAGC 2640
 Db |||||
 2581 GCTCCAGATGTTTCAGTGTGTTTTCAGGGGCTGGAGTTTTTCGTTCTCTCTCTGAGC 2640
 Qy |||||
 2641 CTTTTATCTTTTCTCTCTCTGGAGGAGAAAGTCTATTTTCATGAAGGATGCAATTTTC 2700
 Db |||||
 2641 CTTTTATCTTTTCTCTCTCTGGAGGAGAAAGTCTATTTTCATGAAGGATGCAATTTTC 2700
 Qy |||||
 2701 ATAAAGTCAGCTGTAAATTCAGGCTGTGCATGGGTTCCTTTCAGGAAGGCTTTTAT 2760
 Db |||||
 2701 ATAAAGTCAGCTGTAAATTCAGGCTGTGCATGGGTTCCTTTCAGGAAGGCTTTTAT 2760
 Qy |||||
 2761 TTAATGGAATATAGGAAGCGAGTCACTTCTAGGCGCTTAAATTCACGGAAGAGTGCAC 2820
 Db |||||
 2761 TTAATGGAATATAGGAAGCGAGTCACTTCTAGGCGCTTAAATTCACGGAAGAGTGCAC 2820
 Qy |||||
 2821 TGGAGTCTTTTCTTCTCACTGCTTCTGGGCAACTACTCAGCCCTGTGTGGCTTTA 2880
 Db |||||
 2821 TGGAGTCTTTTCTTCTCACTGCTTCTGGGCAACTACTCAGCCCTGTGTGGCTTTA 2880
 Qy |||||
 2881 TGCAGAGCGTCCGAAACCTTGGAAATCAGGAGACTCGTTCCTTCTGCTTCTGCCATT 2940
 Db |||||
 2881 TGCAGAGCGTCCGAAACCTTGGAAATCAGGAGACTCGTTCCTTCTGCTTCTGCCATT 2940
 Qy |||||
 2941 GGTGGCTGTGCGACCGTGGSCAAGTGTCTCTCTCTCCCTGGGCGCATAGTCTTCTCTCT 3000
 Db |||||
 2941 GGTGGCTGTGCGACCGTGGSCAAGTGTCTCTCTCTCCCTGGGCGCATAGTCTTCTCTCT 3000
 Qy |||||
 3001 ATAAAGACCTTGTGAGCTCTCGTGTCTGTGAACAACATTTCCCTGTGATTTCTCTGAGGG 3060
 Db |||||
 3001 ATAAAGACCTTGTGAGCTCTCGTGTCTGTGAACAACATTTCCCTGTGATTTCTCTGAGGG 3060
 Qy |||||
 3061 GATGTTGAGAGGGAAGGAGGAGCTGAGCAGCTGAGCCACAGGGAGGTGGAGGG 3120
 Db |||||
 3061 GATGTTGAGAGGGAAGGAGGAGCTGAGCAGCTGAGCCACAGGGAGGTGGAGGG 3120
 Qy |||||
 3121 GGACAGGAAGCGCAGGAGAGAGCTGGGTGCTCCATCAGTCTCTCAGTCACTGATCAGTCA 3180
 Db |||||
 3121 GGACAGGAAGCGCAGGAGAGAGCTGGGTGCTCCATCAGTCTCTCAGTCACTGATCAGTCA 3180
 Qy |||||
 3181 CAGGACCGAGAGCCACAAATGCTTTCAGGAAAGCTCAATGAACCCCAAGCCACATTTTCT 3240
 Db |||||
 3181 CAGGACCGAGAGCCACAAATGCTTTCAGGAAAGCTCAATGAACCCCAAGCCACATTTTCT 3240
 Qy |||||
 3241 TCCCTAAGCATAGACATGSCATTTGSCAATAACCAAAAGAAATGAGAGACTTAACCTGGT 3300
 Db |||||
 3241 TCCCTAAGCATAGACATGSCATTTGSCAATAACCAAAAGAAATGAGAGACTTAACCTGGT 3300
 Qy |||||
 3301 GGTAGCTTTTGTGGCTTCAAAAACCTGGGCGCAGAGCAAGTGGAATAATGCCAGATTG 3360
 Db |||||
 3301 GGTAGCTTTTGTGGCTTCAAAAACCTGGGCGCAGAGCAAGTGGAATAATGCCAGATTG 3360
 Qy |||||
 3361 TTAACCTTTTTCACCTGACAGCACCCACGAGCTCAGCAGTGTGCTGACAGCAGG 3420
 Db |||||
 3361 TTAACCTTTTTCACCTGACAGCACCCACGAGCTCAGCAGTGTGCTGACAGCAGG 3420
 Qy |||||
 3421 AGTGACCTGACAGCGCAGGGAGGAGAGAAAGAGAGGGATAGTGTATGAGCAAGAAAG 3480
 Db |||||
 3421 AGTGACCTGACAGCGCAGGGAGGAGAGAAAGAGAGGGATAGTGTATGAGCAAGAAAG 3480
 Qy |||||
 3481 ACAGATTCTTCAAGGCGCAGTGGAAATTTGACACAGGGATTTAGTCCAGTGTATCTG 3540
 Db |||||
 3481 ACAGATTCTTCAAGGCGCAGTGGAAATTTGACACAGGGATTTAGTCCAGTGTATCTG 3540
 Qy |||||
 3541 GTTCTAGGAGCGCAGGCTATATTGTGGGGGAAAAAATCAGTTCAAGGGAGGTGGGAGA 3600
 Db |||||
 3541 GTTCTAGGAGCGCAGGCTATATTGTGGGGGAAAAAATCAGTTCAAGGGAGGTGGGAGA 3600
 Qy |||||
 3601 CCTGATTCTTAATCTATATTTTCTTACAGCTGAGTAAATCTGAGCAAGTCAAG 3660
 Db |||||
 3601 CCTGATTCTTAATCTATATTTTCTTACAGCTGAGTAAATCTGAGCAAGTCAAG 3660
 Qy |||||
 3661 GTAGTAAGCTGAGGCTGTAAGATTACTTAGTTTCTCTCTTATTAGGAACCTCTTTTCTCTGT 3720

Db	4851	GGAAAGAGGAGTATCCAGTTAGCCAAAGTGTCAGGCTGTGCTCTTATTTTATGTA	4910
Qy	181	CAGATGTTGCTCCTGAGAGAGCTATTCTTCAGGAAACATCACATCCAAATATGTAATC	240
Db	4911	CAGATGTTGCTCCTGAGAGAGCTATTCTTCAGGAAACATCACATCCAAATATGTAATC	4970
Qy	241	CATCAACAGGAGCTTAAGAAACAGGAATGAGATGGGCACTTGCCCAAGGAAATGCCAG	300
Db	4971	CATCAACAGGAGCTTAAGAAACAGGAATGAGATGGGCACTTGCCCAAGGAAATGCCAG	5030
Qy	301	GAGAGCAAAATGATGAAAAATAAATTTTCCCTTTGTTTAAATTTTCAAGAAAAATG	360
Db	5031	GAGAGCAAAATGATGAAAAATAAATTTTCCCTTTGTTTAAATTTTCAAGAAAAATG	5090
Qy	361	ATGAGGCCAAATCAATGATAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG	420
Db	5091	ATGAGGCCAAATCAATGATAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG	5150
Qy	421	TAATTAAGTATTTGTTCTCGGGAAGAGACCTCCATGTGAGCTTGTAGGGAAAAATGGAA	480
Db	5151	TAATTAAGTATTTGTTCTCGGGAAGAGACCTCCATGTGAGCTTGTAGGGAAAAATGGAA	5210
Qy	481	AAACGTCAAAAGCATGATCTGATCAGATCCCAAAGTGGATTTATTTAAAAACCCAGAT	540
Db	5211	AAACGTCAAAAGCATGATCTGATCAGATCCCAAAGTGGATTTATTTAAAAACCCAGAT	5270
Qy	541	GGCATCACTCTGGGGAGGCAAGTTTCAGGAAGGTCAATGTAGCAAAAGGACATCAATAAC	600
Db	5271	GGCATCACTCTGGGGAGGCAAGTTTCAGGAAGGTCAATGTAGCAAAAGGACATCAATAAC	5330
Qy	601	AGCAAAATCAAAATTCGCGAAATGCAGAGAGAAATGGGACTGGGAAAGCTTTCATAAC	660
Db	5331	AGCAAAATCAAAATTCGCGAAATGCAGAGAGAAATGGGACTGGGAAAGCTTTCATAAC	5390
Qy	661	AGTGAATAGGCACTTGACCATGTTTCGCAACACCTCCCGTCTATACCAAGGAAACACAAA	720
Db	5391	AGTGAATAGGCACTTGACCATGTTTCGCAACACCTCCCGTCTATACCAAGGAAACACAAA	5450
Qy	721	ATTGACCTGGGCTAGCCCTGACCTTTCAAGGAAATNTGAAAACTGAGAGCAAAACAAA	780
Db	5451	ATTGACCTGGGCTAGCCCTGACCTTTCAAGGAAATNTGAAAACTGAGAGCAAAACAAA	5510
Qy	781	GACATGTTTAAAGGCAACAGAACATTTGTGAGCTTCAAGAGCAGAGTCCCTCAGCA	840
Db	5511	GACATGTTTAAAGGCAACAGAACATTTGTGAGCTTCAAGAGCAGAGTCCCTCAGCA	5570
Qy	841	GGGACCTGAGGCAATTTGCCCTTTAGGAAGGCAGTCTTTTAAAGGAATCTTAAAGAACTC	900
Db	5571	GGGACCTGAGGCAATTTGCCCTTTAGGAAGGCAGTCTTTTAAAGGAATCTTAAAGAACTC	5630
Qy	901	TTGAAAGATCATGAATTTTAAACATTTTAAAGTATAAAACAAATATGCGATGCAATACAG	960
Db	5631	TTGAAAGATCATGAATTTTAAACATTTTAAAGTATAAAACAAATATGCGATGCAATACAG	5690
Qy	961	TTTATAGATGGTCCCAATTTTAAAGTACAGGATACAGGATACAGTCTCCAGCTCC	1020
Db	5691	TTTATAGATGGTCCCAATTTTAAAGTACAGGATACAGGATACAGTCTCCAGCTCC	5750
Qy	1021	GGATAGTCAAGAAATCACTGTGTGCCCATCTTAACCTTTTTCAGAAATGATC	1080
Db	5751	GGATAGTCAAGAAATCACTGTGTGCCCATCTTAACCTTTTTCAGAAATGATC	5810
Qy	1081	TGTATAGCCCTCACAACAGGCGCGATGTGTGACCTTCAACCATCATCAACCCAA	1140
Db	5811	TGTATAGCCCTCACAACAGGCGCGATGTGTGACCTTCAACCATCATCAACCCAA	5870
Qy	1141	GTCCCTCAACATTTGTTAAAGTGTATCTCAGTAGGTCCCATTTACAAATGCCACCTCCC	1200
Db	5871	GTCCCTCAACATTTGTTAAAGTGTATCTCAGTAGGTCCCATTTACAAATGCCACCTCCC	5930
Qy	1201	TGTGACGCCCATCCCGCTCCACAGGAAGTCTCCCTCCTCAGTGTCTGTGATCAGGATGT	1260
Db	5931	TGTGACGCCCATCCCGCTCCACAGGAAGTCTCCCTCCTCAGTGTCTGTGATCAGGATGT	5990
Qy	1261	TACAGCCAGAGCTCCCGTGAGGGTGAGGGTCTGTGTCTTACACCTACTGTATGCTCTAC	1320
Db	5991	TACAGCCAGAGCTCCCGTGAGGGTGAGGGTCTGTGTCTTACACCTACTGTATGCTCTAC	6050
Qy	1321	ACCTGAGCTCACTGCACAACTCTGCTCCCAAGTTCAGGCAATTTCTCTGTCTCAGCCTCC	1380
Db	6051	ACCTGAGCTCACTGCACAACTCTGCTCCCAAGTTCAGGCAATTTCTCTGTCTCAGCCTCC	6110
Qy	1381	CGGTAGCTGGGACTACAGGGCGACGCGCGCTAAATTTTGTATTTGTAGTAGAGTGG	1440
Db	6111	CGGTAGCTGGGACTACAGGGCGACGCGCGCTAAATTTTGTATTTGTAGTAGAGTGG	6170
Qy	1441	GTCTTACCATAATTTAGCCGCTGTGAACTCTGACCTCAGGTGATCCACCACTC	1500
Db	6171	GTCTTACCATAATTTAGCCGCTGTGAACTCTGACCTCAGGTGATCCACCACTC	6230
Qy	1501	AGCTCTCTAAAGTGTGGGATTTACAGGCATGAGTCAACGCGCGCGGCAAGGGTCAAGTGT	1560
Db	6231	AGCTCTCTAAAGTGTGGGATTTACAGGCATGAGTCAACGCGCGCGGCAAGGGTCAAGTGT	6290
Qy	1561	TTAATAAGGAATTAATTTGAATGTTTCTAAACCAACAGGGAACAGACAAAGCTGTGA	1620
Db	6291	TTAATAAGGAATTAATTTGAATGTTTCTAAACCAACAGGGAACAGACAAAGCTGTGA	6350
Qy	1621	TAATTTTACGGGATTTCTGGGATGGGAAATGGTGCATGAGCTGCCTAGTCCCAGAC	1680
Db	6351	TAATTTTACGGGATTTCTGGGATGGGAAATGGTGCATGAGCTGCCTAGTCCCAGAC	6410
Qy	1681	CACGTGCTCTCATCATCTTCTCCCTCATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	1740
Db	6411	CACGTGCTCTCATCATCTTCTCCCTCATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	6470
Qy	1741	CACCATGCTTTTGTGGTAAAGCTTCCACATCTGTAATAAGAGTATACATAAAGCTAG	1800
Db	6471	CACCATGCTTTTGTGGTAAAGCTTCCACATCTGTAATAAGAGTATACATAAAGCTAG	6530
Qy	1801	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGGAGGAGGAGGAGGAGGAGGAGGAGG	1860
Db	6531	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGGAGGAGGAGGAGGAGGAGGAGGAGG	6590
Qy	1861	TGAAGCCCGCGGAGAGGTTTCTCTCTCAGCTGGGGAGGAGGAGGAGGAGGAGGAGG	1920
Db	6591	TGAAGCCCGCGGAGAGGTTTCTCTCTCAGCTGGGGAGGAGGAGGAGGAGGAGGAGG	6650
Qy	1921	TGGGTGTCTGAGCAACCTGCCAGCCGCTGCCACTGGTGTGTTTGTATCACTCTCTAGG	1980
Db	6651	TGGGTGTCTGAGCAACCTGCCAGCCGCTGCCACTGGTGTGTTTGTATCACTCTCTAGG	6710
Qy	1981	GACCTGTGCTTCTATTTCTGTGTGACTGTGTTTCTATCCAGGCAATTCATTTGACAAAT	2040
Db	6711	GACCTGTGCTTCTATTTCTGTGTGACTGTGTTTCTATCCAGGCAATTCATTTGACAAAT	6770
Qy	2041	TATTGAGTACTTATATCTGCCAGACACAGAGCAAAATGGTGAGCAAAATTAATAGCCA	2100
Db	6771	TATTGAGTACTTATATCTGCCAGACACAGAGCAAAATGGTGAGCAAAATTAATAGCCA	6830
Qy	2101	CCTACCTCTGAGGAGTGAAGTTCTCATGGAAGAGGAGGAGGAGGAGGAGGAGGAGG	2160
Db	6831	CCTACCTCTGAGGAGTGAAGTTCTCATGGAAGAGGAGGAGGAGGAGGAGGAGGAGG	6890
Qy	2161	GCCAACTTAAACCCAGTGTGAAAGAAAGGAAATTAACCATCTTTGAAGAAATTTGCGC	2220
Db	6891	GCCAACTTAAACCCAGTGTGAAAGAAAGGAAATTAACCATCTTTGAAGAAATTTGCGC	6950
Qy	2221	AGCATCTTAAACAGGCGCACTCCCTAGCGCCCTGCTGCTCCTCCTGCTGCTGCTGCTG	2280
Db	6951	AGCATCTTAAACAGGCGCACTCCCTAGCGCCCTGCTGCTCCTCCTGCTGCTGCTGCTG	7010
Qy	2281	CCCCAAGCCCGAGTCTTCAAGCCCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	2340
Db	7011	CCCCAAGCCCGAGTCTTCAAGCCCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	7070

Qy	2341	GCCTCGCTTCCCGTGAATCGTCTGTGTGCATCTGAGCTGGAGACTCTCTTGGCTCCAGGCT	2400
Db	7071	GCCTCGCTTCCCGTGAATCGTCTGTGTGCATCTGAGCTGGAGACTCTCTTGGCTCCAGGCT	7130
Qy	2401	CCAGAAAGGAAATGGAGAGGAAACTTAGTCTTAACGGAGAACTCTGAGAGGGACAGTGTTC	2460
Db	7131	CCAGAAAGGAAATGGAGAGGAAACTTAGTCTTAACGGAGAACTCTGAGAGGGACAGTGTTC	7190
Qy	2461	CTCAGAGGAAAGGGGGCTCCACGTCCAGGAGAAATTCACGAGGTGGGACTCGACGGAG	2520
Db	7191	CTCAGAGGAAAGGGGGCTCCACGTCCAGGAGAAATTCACGAGGTGGGACTCGACGGAG	7250
Qy	2521	TGGGACGCTGGGGCTGAGCGGTGTGAAAGGACGGAAGGTGAAAGGGCAAGCGTGAA	2580
Db	7251	TGGGACGCTGGGGCTGAGCGGTGTGAAAGGACGGAAGGTGAAAGGGCAAGCGTGAA	7310
Qy	2581	GCTGCCACAGATGTTACAGTGTTGTTCACGGGGCTGGGAGTTTCCGTTCCTCTGTGAGC	2640
Db	7311	GCTGCCACAGATGTTACAGTGTTGTTCACGGGGCTGGGAGTTTCCGTTCCTCTGTGAGC	7370
Qy	2641	CTTTTATCTTTTCTCTGCTTGGAGGAGAAAGTCTATTTTCAATGAAGGATGCAGTTTC	2700
Db	7371	CTTTTATCTTTTCTCTGCTTGGAGGAGAAAGTCTATTTTCAATGAAGGATGCAGTTTC	7430
Qy	2701	ATAAAGTCAGCTGTTAAATTCACGGGTGTGCATGGGTTTTCTTCAAGAGCCCTTAT	2760
Db	7431	ATAAAGTCAGCTGTTAAATTCACGGGTGTGCATGGGTTTTCTTCAAGAGCCCTTAT	7490
Qy	2761	TTAATGGGAATATAGGAAGCGAGCTCAATTTCTTAGCCGTTAATTCACGGAAGAGTGAC	2820
Db	7491	TTAATGGGAATATAGGAAGCGAGCTCAATTTCTTAGCCGTTAATTCACGGAAGAGTGAC	7550
Qy	2821	TGGAGTCTTTTCTTTCAATGCTTCTGCGGCAACTACTCAGCCCTGTGTGAGCTTGGCTTA	2880
Db	7551	TGGAGTCTTTTCTTTCAATGCTTCTGCGGCAACTACTCAGCCCTGTGTGAGCTTGGCTTA	7610
Qy	2881	TGCAAGACGCTGCGAAACCTTGTGAATCAGAGACTCGGTTTTCTTCTGGTCTGCATTT	2940
Db	7611	TGCAAGACGCTGCGAAACCTTGTGAATCAGAGACTCGGTTTTCTTCTGGTCTGCATTT	7670
Qy	2941	GGTTGGCTGTGCGACCGTGGGCAAGTGCTCTCCTCCCTGGGCCATAGTCTTCTCTGCT	3000
Db	7671	GGTTGGCTGTGCGACCGTGGGCAAGTGCTCTCCTCCCTGGGCCATAGTCTTCTCTGCT	7730
Qy	3001	ATAAAGACCCCTTGACGCTCTCGTGTCTGTGAAACACTTCCCTGTGATTTCTCTGAGGGG	3060
Db	7731	ATAAAGACCCCTTGACGCTCTCGTGTCTGTGAAACACTTCCCTGTGATTTCTCTGAGGGG	7790
Qy	3061	GGATGTTGAGAGGGGAAGGAGCGACAGCTGGAGCAGTGAAGCCACAGGGGAGGTGAGGG	3120
Db	7791	GGATGTTGAGAGGGGAAGGAGCGACAGCTGGAGCAGTGAAGCCACAGGGGAGGTGAGGG	7850
Qy	3121	GGACAGGAAGCGCAGAGAGCTGGGTGCTCCATCAGTCTCTCACTGATCAGTCCAGATC	3180
Db	7851	GGACAGGAAGCGCAGAGAGCTGGGTGCTCCATCAGTCTCTCACTGATCAGTCCAGATC	7910
Qy	3181	CAGGACCAGAGCCACAAATGCTTCAGGAAAGCTCAATGAACCCAAAGCCACATTTTCT	3240
Db	7911	CAGGACCAGAGCCACAAATGCTTCAGGAAAGCTCAATGAACCCAAAGCCACATTTTCT	7970
Qy	3241	TCCCTAAGCATPAGCAATGGCATTTGCCAATAACCAAAAGAAATGACAGACTAATCTGCT	3300
Db	7971	TCCCTAAGCATPAGCAATGGCATTTGCCAATAACCAAAAGAAATGACAGACTAATCTGCT	8030
Qy	3301	GGTAGCTTTTGGCTGGCATTTCAAAAATGGGCCAGAGCAAGTGGAAATGCCAGAGATTG	3360
Db	8031	GGTAGCTTTTGGCTGGCATTTCAAAAATGGGCCAGAGCAAGTGGAAATGCCAGAGATTG	8090
Qy	3361	TTAAACTTTTTCACCTGACAGCACCCACGACGCTCAGCAGTGTGCTGTGACAGACCGG	3420
Db	8091	TTAAACTTTTTCACCTGACAGCACCCACGACGCTCAGCAGTGTGCTGTGACAGACCGG	8150
Qy	3421	AGTGACCTTGACCGCAGGGGAGGAGAAAGAGAGGGGATAGTGTATGACGCAAGAAAG	3480

Db	8151	AGTGA	CCTG	CGG	CAGGG	GAGG	AGN	AGAAA	AGAG	GGG	ATAGT	GTATG	AGCA	AGAA	G	8210	
Qy	3481	ACAGAT	TCATT	CA	AGG	CAGT	GGG	AA	TTGA	CCAC	AGGG	AA	TATAGT	CCAC	GTGAT	CCTGG	3540
Db	8211	ACAGAT	TCATT	CA	AGG	CAGT	GGG	AA	TTGA	CCAC	AGGG	AA	TATAGT	CCAC	GTGAT	CCTGG	8270
Qy	3541	GTTC	TAGG	AGG	CAGG	CGCT	ATAT	TGTGG	GGG	AAAA	AAAT	TCAGT	TC	AGG	GAAGT	CCGGG	3600
Db	8271	GTTC	TAGG	AGG	CAGG	CGCT	ATAT	TGTGG	GGG	AAAA	AAAT	CAGT	TC	AGG	GAAGT	CCGGG	8330
Qy	3601	CCTG	ATTC	TT	CT	AA	TACT	ATAT	TTTT	CC	TTT	CA	AGCT	GTAG	CTAA	TTCTG	3660
Db	8331	CCTG	ATTC	TT	CT	AA	TACT	ATAT	TTTT	CC	TTT	CA	AGCT	GTAG	CTAA	TTCTG	8390
Qy	3661	GTAG	TAACT	GAG	CGT	GT	AA	GAT	TACT	AGT	TTCT	CC	TAT	TAG	GAAT	CTCTTT	3720
Db	8391	GTAG	TAACT	GAG	CGT	GT	AA	GAT	TACT	AGT	TTCT	CC	TAT	TAG	GAAT	CTCTTT	8450
Qy	3721	GGAG	TTAG	CAG	CAC	AGG	CGCA	ATCC	CGT	TTCT	TTT	AA	CAGG	AA	GAAT	CA	3780
Db	8451	GGAG	TTAG	CAG	CAC	AGG	CGCA	ATCC	CGT	TTCT	TTT	AA	CAGG	AA	GAAT	CA	8510
Qy	3781	TAAAG	CCAA	CAG	AT	CAAG	CC	TAGT	CT	TGCT	GA	CTAT	AT	GAT	TGGT	TTTT	3840
Db	8511	TAAAG	CCAA	CAG	AT	CAAG	CC	TAGT	CT	TGCT	GA	CTAT	AT	GAT	TGGT	TTTT	8570
Qy	3841	CATT	TCAG	CGAT	GT	TT	ACT	CTGA	TT	CA	GA	AAAT	TGAG	ACT	AGT	ACCC	3900
Db	8571	CATT	TCAG	CGAT	GT	TT	ACT	CTGA	TT	CA	GA	AAAT	TGAG	ACT	AGT	ACCC	8630
Qy	3901	TAA	ACAA	CA	CC	CAGT	GT	AAAT	TC	CAAGT	TC	AGG	CTT	AA	TCG	CAG	3959
Db	8631	TAA	ACAA	CA	CC	CAGT	GT	AAAT	TC	CAAGT	TC	AGG	CTT	AA	TCG	CAG	8690
Qy	3960	AAGA	AT	TAGA	AT	CT	TT	TAG	CAAA	CT	CTG	TTT	CT	CA	CT	CG	4019
Db	8691	AAGA	AT	TAGA	AT	CT	TT	TAG	CAAA	CT	CTG	TTT	CT	CA	CT	CG	8749
Qy	4020	CAGT	TTT	GG	AA	TAT	TT	ACT	TC	CA	AGT	AT	GTG	AT	TT	AA	4079
Db	8750	CAGT	TTT	GG	AA	TAT	TT	ACT	TC	CA	AGT	AT	GTG	AT	TT	AA	8809
Qy	4080	TTG	CT	CA	AGG	CAAT	CA	TAT	TT	CA	AGT	GGC	TT	AA	AGT	TT	4139
Db	8810	TTG	CT	CA	AGG	CAAT	CA	TAT	TT	CA	AGT	GGC	TT	AA	AGT	TT	8869
Qy	4140	TTT	AT	T	GGC	TAT	T	GGC	AT	T	T	T	T	T	T	T	4199
Db	8870	TTT	AT	T	GGC	TAT	T	GGC	AT	T	T	T	T	T	T	T	8929
Qy	4200	GGG	AT	T	AA	CCT	CA	CAGT	CC	CA	GA	AGC	CT	GT	GA	AT	4259
Db	8930	GGG	AT	T	AA	CCT	CA	CAGT	CC	CA	GA	AGC	CT	GT	GA	AT	8989
Qy	4260	TTG	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	4319
Db	8990	TTG	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	9049
Qy	4320	ACT	CA	AGT	CGG	TAAT	TA	AC	AGT	AC	CT	GT	AT	T	T	T	4379
Db	9050	ACT	CA	AGT	CGG	TAAT	TA	AC	AGT	AC	CT	GT	AT	T	T	T	9109
Qy	4380	TTT	ATA	CT	AT	T	TA	CT	AGT	TG	TT	CG	AT	GT	TAAGT	GAA	4439
Db	9110	TTT	ATA	CT	AT	T	TA	CT	AGT	TG	TT	CG	AT	GT	TAAGT	GAA	

QY 961 TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAGGATAAGTGTCCAGCTCC 1020
DB 961 TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAGGATAAGTGTCCAGCTCC 1020
QY 1021 GGATAGTTCAGAAATCATTTAGAAATCACTGTGTCCCATCTTAACATTTTTCAGAAATGATC 1080
DB 1021 GGATAGTTCAGAAATCATTTAGAAATCACTGTGTCCCATCTTAACATTTTTCAGAAATGATC 1080
QY 1081 TGTATAGCCCTCACACAGGCCCGATGTGTCTGACCTTACACCACTCTACACCCAA 1140
DB 1081 TGTATAGCCCTCACACAGGCCCGATGTGTCTGACCTTACACCACTCTACACCCAA 1140
QY 1141 GTGCTCAACCATTTTAAAGTGTCTATCTCAGTAGGTCCCATTTACAAATGCCACCTCC 1200
DB 1141 GTGCTCAACCATTTTAAAGTGTCTATCTCAGTAGGTCCCATTTACAAATGCCACCTCC 1200
QY 1201 TGTGACGCCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCATCAGATGT 1260
DB 1201 TGTGACGCCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCATCAGATGT 1260
QY 1261 TACAGCCAGAAAGTCTCCGTGAGGTGAGGTCTGTGTCTTACACTTACCTTGTATCTCTAC 1320
DB 1261 TACAGCCAGAAAGTCTCCGTGAGGTGAGGTCTGTGTCTTACACTTACCTTGTATCTCTAC 1320
QY 1321 ACTGAGCTCACTGCAACTCTGCTCCAGAGTTCAAGCAATTTCTCTGTCTCAGCTCC 1380
DB 1321 ACTGAGCTCACTGCAACTCTGCTCCAGAGTTCAAGCAATTTCTCTGTCTCAGCTCC 1380
QY 1381 CGGTAGCTGGGACTACAGCGCACCGCCGGCTAAATTTTGTATTTAGTAGAGATGG 1440
DB 1381 CGGTAGCTGGGACTACAGCGCACCGCCGGCTAAATTTTGTATTTAGTAGAGATGG 1440
QY 1441 GTTTACCAATTTAGCCCGGTCTTGAATCTCTGAACTCTGAACTCTGAACTCTGAACTCTG 1500
DB 1441 GTTTACCAATTTAGCCCGGTCTTGAATCTCTGAACTCTGAACTCTGAACTCTGAACTCTG 1500
QY 1501 AGCTCTTAAAGTCTGGGATTTACAGCATGAGTCAAGCGCCCGCCAGGCTCAGTGT 1560
DB 1501 AGCTCTTAAAGTCTGGGATTTACAGCATGAGTCAAGCGCCCGCCAGGCTCAGTGT 1560
QY 1561 TTAATAAGGAATAAATGAAATGTTTAACTAAACCAACAGGAAACAGACAAAGCTGTA 1620
DB 1561 TTAATAAGGAATAAATGAAATGTTTAACTAAACCAACAGGAAACAGACAAAGCTGTA 1620
QY 1621 TAAATTCAGGATTTCTGGATGGGAAATGGTCCATGAGTCCCTGCTGCTAGTCCAGAC 1680
DB 1621 TAAATTCAGGATTTCTGGATGGGAAATGGTCCATGAGTCCCTGCTGCTAGTCCAGAC 1680
QY 1681 CACTGCTCTCATCATCTTCTCCCTCATCTCTCATTTTCAAGGCTAAGTTTACATTTTAT 1740
DB 1681 CACTGCTCTCATCATCTTCTCCCTCATCTCTCATTTTCAAGGCTAAGTTTACATTTTAT 1740
QY 1741 CACATGCTTTTGTGGTAAAGCTCCACATCGTTTACTGAAATAGAGTATACATAAAGTAG 1800
DB 1741 CACATGCTTTTGTGGTAAAGCTCCACATCGTTTACTGAAATAGAGTATACATAAAGTAG 1800
QY 1801 TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGGAGGCGATATCCCGAGACTCT 1860
DB 1801 TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGGAGGCGATATCCCGAGACTCT 1860
QY 1861 TGAAGCCCGGAGAGGTTTCTCTCCAGCTGGGAGCCCTGCAAGCACCCGGGCTCC 1920
DB 1861 TGAAGCCCGGAGAGGTTTCTCTCCAGCTGGGAGCCCTGCAAGCACCCGGGCTCC 1920
QY 1921 TGGGTGTCTGAGCAACTGCTCCAGCCCGTGCCTGCTGTTTGTATCACTCTCTAGG 1980
DB 1921 TGGGTGTCTGAGCAACTGCTCCAGCCCGTGCCTGCTGTTTGTATCACTCTCTAGG 1980
QY 1981 GACCTGTGTCTTCTATTTCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 2040
DB 1981 GACCTGTGTCTTCTATTTCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 2040
QY 2041 TATTGAGTACTTATATCTGCCAGACACAGGACAAATGGTGTAGCAAGCAGTCACTGC 2100

DB 2041 TATTGAGTACTTATATCTGCCAGACACAGGACAAATGGTGTAGCAAGCAGTCACTGC 2100
QY 2101 CCTACCTTCGTGGAGTGCAGTGTCTCATGGAAGACGTGAGAAAGAAATTAATAGCCA 2160
DB 2101 CCTACCTTCGTGGAGTGCAGTGTCTCATGGAAGACGTGAGAAAGAAATTAATAGCCA 2160
QY 2161 GCCAACTTAAACCCAGTCTGAAAGAAAGAAATTAACACCATCTTGAAGAAATTTGGGC 2220
DB 2161 GCCAACTTAAACCCAGTCTGAAAGAAAGAAATTAACACCATCTTGAAGAAATTTGGGC 2220
QY 2221 AGCATCCCTTAAACAGGCCACTCTCCTAGCGCCCTCTGCTCCATCTGTCGCCGAGG 2280
DB 2221 AGCATCCCTTAAACAGGCCACTCTCCTAGCGCCCTCTGCTCCATCTGTCGCCGAGG 2280
QY 2281 CCCCAAGCCGAGTCTTCCAAAGCTCTCTCTCATAGTCAAGCTCTGAGCTGAGCTG 2340
DB 2281 CCCCAAGCCGAGTCTTCCAAAGCTCTCTCTCATAGTCAAGCTCTGAGCTGAGCTG 2340
QY 2341 GCCTCGCTTCCCGTGAATCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2400
DB 2341 GCCTCGCTTCCCGTGAATCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2400
QY 2401 CCAGAAAGAAATGGAGAGGAACTAGTCTAAACGGAATCTGAGGGGACAGTGTTC 2460
DB 2401 CCAGAAAGAAATGGAGAGGAACTAGTCTAAACGGAATCTGAGGGGACAGTGTTC 2460
QY 2461 CTGAGAGGAAAGGGCCCTCCAGTCTCAGAGGAATTCAGAGGAGTGGGACATGAGGAG 2520
DB 2461 CTGAGAGGAAAGGGCCCTCCAGTCTCAGAGGAATTCAGAGGAGTGGGACATGAGGAG 2520
QY 2521 TGGGAGCCCTGGGCTGAGCGGTGCTGAAAGGAGGAAGTGAAGGAGGAGGAGTGA 2580
DB 2521 TGGGAGCCCTGGGCTGAGCGGTGCTGAAAGGAGGAAGTGAAGGAGGAGGAGTGA 2580
QY 2581 GCTGCCAGAGTGTTCAGTGTGTTTCAAGGGGCTGGAGTGTTCCTGCTCTGAGC 2640
DB 2581 GCTGCCAGAGTGTTCAGTGTGTTTCAAGGGGCTGGAGTGTTCCTGCTCTGAGC 2640
QY 2641 CTTTTTATCTTTCTCTGCTTGGAGGAGAAAGTCTATTTTCAAGGAGTGGAGTTC 2700
DB 2641 CTTTTTATCTTTCTCTGCTTGGAGGAGAAAGTCTATTTTCAAGGAGTGGAGTTC 2700
QY 2701 ATAAAGTCAGCTGTTTAAATTTCCAGGGTGTGATCGGTTTCTTTCACGAGGCTTTAT 2760
DB 2701 ATAAAGTCAGCTGTTTAAATTTCCAGGGTGTGATCGGTTTCTTTCACGAGGCTTTAT 2760
QY 2761 TTAATGGGAATATAGGAAGCAGCTCATTTTCTAGGCCGTAAATTCACGGAAGAGTGAC 2820
DB 2761 TTAATGGGAATATAGGAAGCAGCTCATTTTCTAGGCCGTAAATTCACGGAAGAGTGAC 2820
QY 2821 TGGAGTCTTTTCTTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTT 2880
DB 2821 TGGAGTCTTTTCTTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTT 2880
QY 2881 TGCAAGCGTTCGAAACCTTTGGAATCAGGAGCTCGGTTTCTTTCTTTCTTTCTTTCTTT 2940
DB 2881 TGCAAGCGTTCGAAACCTTTGGAATCAGGAGCTCGGTTTCTTTCTTTCTTTCTTTCTTT 2940
QY 2941 GGTTCGCTGTGCGACCGTGGGCAAGTGTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCT 3000
DB 2941 GGTTCGCTGTGCGACCGTGGGCAAGTGTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCT 3000
QY 3001 ATAAAGACCTTTCAGCTCTGTGTCTGTGAACTTCTTCTTCTTCTTCTTCTTCTTCTTCT 3060
DB 3001 ATAAAGACCTTTCAGCTCTGTGTCTGTGAACTTCTTCTTCTTCTTCTTCTTCTTCTTCT 3060
QY 3061 GGATGTTGAGAGGGAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3120
DB 3061 GGATGTTGAGAGGGAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3120
QY 3121 GGACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3180


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; Publication No. US20050170353A1
; GENERAL INFORMATION:
; APPLICANT: Syntex Corporation
; APPLICANT: ASANO, Kaoru
; APPLICANT: TAKAHATA, Takayuki
; APPLICANT: NUMADA, Shigehiro
; APPLICANT: MASAGO, Akinoiri
; APPLICANT: KOUCHI, Yasuhiro
; TITLE OF INVENTION: GENE EXAMINATION METHOD FOR JUDGING THE ONSET RISK OF GLAUCOMA
; FILE REFERENCE: 083447
; CURRENT APPLICATION NUMBER: US/10/509,595
; CURRENT FILING DATE: 2004-09-29
; PRIOR APPLICATION NUMBER: JP 2002-093443
; PRIOR FILING DATE: 2002-03-29
; PRIOR APPLICATION NUMBER: PCT/JP03/03307
; PRIOR FILING DATE: 2003-03-19
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 1
; LENGTH: 6000
; TYPE: DNA
; ORGANISM: Homo sapiens
; US-10-509-595-1

Query Match      76.2%; Score 4017.4; DB 9; Length 6000;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 4052; Conservative 0; Mismatches 1; Indels 4; Gaps 3;

QY 1216 GCTCCACAGGAAGTCTCCCACTAGACTCTGCACTGATGCTTACACCTACCTGATGCTTACACCTGAGTCACTGC 1335
DB 1 GCTCCACAGGAAGTCTCCCACTAGACTCTGCACTGATGCTTACACCTACCTGATGCTTACACCTGAGTCACTGC 1275

QY 1276 CGTGAGGCTGAGGCTGTGCTCTTACACCTACCTGATGCTTACACCTGATGCTTACACCTGAGTCACTGC 1335
DB 61 CGTGAGGCTGAGGCTGTGCTCTTACACCTACCTGATGCTTACACCTGAGTCACTGC 120

QY 1336 AACCTCTGCTCCAGGATCAAGCAATCTCTGCTGATGCTTACACCTGATGCTTACACCTGAGTCACTGC 1395
DB 121 AACCTCTGCTCCAGGATCAAGCAATCTCTGCTGATGCTTACACCTGATGCTTACACCTGAGTCACTGC 180

QY 1396 ACAGGCGCAGCCGGCTAATTTTGTATTGTAGTAGATGAGTGGGTTTCCACCATATTAG 1455
DB 181 ACAGGCGCAGCCGGCTAATTTTGTATTGTAGTAGATGAGTGGGTTTCCACCATATTAG 240

QY 1456 CCGCGCTGCTTGAACCTCTGACCTCAGGTGATGCCACCACTCAGCTCCTTAAAGTGC 1515
DB 241 CCGCGCTGCTTGAACCTCTGACCTCAGGTGATGCCACCACTCAGCTCCTTAAAGTGC 300

QY 1516 TGGGATTAAGGATAGTCAACCGCGCCCGCCGAGGTCAGTGTGTTAATAAGGAATAC 1575
DB 301 TGGGATTAAGGATAGTCAACCGCGCCCGCCGAGGTCAGTGTGTTAATAAGGAATAC 360

QY 1576 TTGAAATGTTTACTTAACCAAGGAAACAGACAAAGCTGTGATAATTTTCCAGGATTC 1635
DB 361 TTGAAATGTTTACTTAACCAAGGAAACAGACAAAGCTGTGATAATTTTCCAGGATTC 420

QY 1636 TTGGGATGGGAATGGTGCATGAGTCCCTGCTGCTAGTCCAGACCACTGCTCCTCATCA 1695
DB 421 TTGGGATGGGAATGGTGCATGAGTCCCTGCTGCTAGTCCAGACCACTGCTCCTCATCA 480

QY 1696 CTTTCTTCCCTCATFCCTCATTTTTCAGGCTAAGTTACATTTTATTCACCATGCTTTTGTG 1755
DB 481 CTTTCTTCCCTCATFCCTCATTTTTCAGGCTAAGTTACATTTTATTCACCATGCTTTTGTG 540

QY 1756 GTAAGCTCCACATCGTTACTGAATAAGAGTATACATAAAGTGTGATTTTCCATTTTGGGCCCA 1815
DB 541 GTAAGCTCCACATCGTTACTGAATAAGAGTATACATAAAGTGTGATTTTCCATTTTGGGCCCA 600

QY 1816 TCTGTGTGTGTATAGGGAGGAGGCGATACCCAGAGACTCCTTGAAGCCCCCGGCGAG 1875
DB 601 TCTGTGTGTGTATAGGGAGGAGGCGATACCCAGAGACTCCTTGAAGCCCCCGGCGAG 660

QY 1876 AGGTTTCTCTCCAGCTGGGGGAGCCCTGCAAGCACCCCGGGTCTCTGGGTGCTCTGAGCA 1935

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DB 661 AGGTTTCTCTCCAGCTGGGGGAGCCCTGCAAGCAACCCGGGTCTCTGGGTGCTCTGAGCA 720
QY 1936 ACCTGCCAGCCCGTGCACACTGGTGTGTTTGTATCATCTCTTAGGAGACTGTGTCTTCT 1995
DB 721 ACCTGCCAGCCCGTGCACACTGGTGTGTTTGTATCATCTCTTAGGAGACTGTGTCTTCT 780
QY 1996 ATTTCTGTGTGACTCGTTCATTCATCCAGGCAATTCATTGACAAATTTATTGAGTACTATA 2055
DB 781 ATTTCTGTGTGACTCGTTCATTCATCCAGGCAATTCATTGACAAATTTATTGAGTACTATA 840
QY 2056 TCTGCCAGACACACAGAGACAAATGCTGAGCAAAAGCAGTCACTGCGCTACCTTCTGTTGAG 2115
DB 841 TCTGCCAGACACACAGAGACAAATGCTGAGCAAAAGCAGTCACTGCGCTACCTTCTGTTGAG 900
QY 2116 GTGACAGTTTCTCATGGAAGACGTGCAGAAAGAAATTAATAGCCAGCCAACTTAAACCCA 2175
DB 901 GTGACAGTTTCTCATGGAAGACGTGCAGAAAGAAATTAATAGCCAGCCAACTTAAACCCA 960
QY 2176 GTGCTGAAAGAAAGAAATTAACACCATCTTGAAGAAATTTGTGGGAGCATCCCTTAACAA 2235
DB 961 GTGCTGAAAGAAAGAAATTAACACCATCTTGAAGAAATTTGTGGGAGCATCCCTTAACAA 1020
QY 2236 GGCCACCTCCCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGGCCCCCAAGCCGAGT 2295
DB 1021 GGCCACCTCCCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGGCCCCCAAGCCGAGT 1080
QY 2296 CTTTCAAGCTCTCTCTCCATCAGTCAAGCCCTGAGCTGCGCTGCTGCTTCCCGTG 2355
DB 1081 CTTTCAAGCTCTCTCTCCATCAGTCAAGCCCTGAGCTGCGCTGCTGCTTCCCGTG 1140
QY 2356 AATCGTCTCTGCTGATCTGAGCTGGAGACTCCTTGGCTCAGGCTCCAGAAAGGAATGG 2415
DB 1141 AATCGTCTCTGCTGATCTGAGCTGGAGACTCCTTGGCTCAGGCTCCAGAAAGGAATGG 1200
QY 2416 ACAGGGAAGTACTAGTCTAACGGAGAACTCTGAGGGGACAGTGTTCCTCAGAGGAAAGGG 2475
DB 1201 ACAGGGAAGTACTAGTCTAACGGAGAACTCTGAGGGGACAGTGTTCCTCAGAGGAAAGGG 1260
QY 2476 GCCTCCACCTCAGGAGAAATTCAGAGGTGGGACTGACGAGGTGGGAGCGCTGGGC 2535
DB 1261 GCCTCCACCTCAGGAGAAATTCAGAGGTGGGACTGACGAGGTGGGAGCGCTGGGC 1320
QY 2536 TGAGCGGTGCTGAAAGGAGGAGGTAAGAGGCGAGGCTGAGCTGCCAGAGTTC 2595
DB 1321 TGAGCGGTGCTGAAAGGAGGAGGTAAGAGGCGAGGCTGAGCTGCCAGAGTTC 1380
QY 2596 AGTGTGTTTTCAGCGGCTGGGAGTTTCCGTTGCTTCTGCTGAGCCCTTTTATCTTTCT 2655
DB 1381 AGTGTGTTTTCAGCGGCTGGGAGTTTCCGTTGCTTCTGCTGAGCCCTTTTATCTTTCT 1440
QY 2656 CTGCTTGGAGGAGAAAGTCTATTTCATGAAGGAGTGCAGTTTCATAAGTCAAGTGT 2715
DB 1441 CTGCTTGGAGGAGAAAGTCTATTTCATGAAGGAGTGCAGTTTCATAAGTCAAGTGT 1500
QY 2716 AAAATTCAGGCTGTCATGGGTTTCTTCCAGAGGCTTTTATTAATGGAATATAG 2775
DB 1501 AAAATTCAGGCTGTCATGGGTTTCTTCCAGAGGCTTTTATTAATGGAATATAG 1560
QY 2776 GAAGCAGCTCATTTTCTTAGGCGGTAAATTCACGGAAGAGTGCAGTGGAGTCTTTCTTT 2835
DB 1561 GAAGCAGCTCATTTTCTTAGGCGGTAAATTCACGGAAGAGTGCAGTGGAGTCTTTCTTT 1620
QY 2836 CATGCTCTCTGGGCAACTACTCAGCCCTGCTGGTGTGAGTCTGGCTTATGCAAGCGTCA 2895
DB 1621 CATGCTCTCTGGGCAACTACTCAGCCCTGCTGGTGTGAGTCTGGCTTATGCAAGCGTCA 1680
QY 2896 AACCTTGAATCAGGAGACTCGGTTTCTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 2955
DB 1681 AACCTTGAATCAGGAGACTCGGTTTCTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1740
QY 2956 CGTGGGCAAGTGTCTCTCTTCCCTGGGCAATAGTCTTCTGCTATAAGACCTTGCA 3015

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Db	1741	C	T	G	G	G	A	A	G	T	G	T	C	T	C	T	C	C	T	G	G	G	C	A	T	A	G	T	C	T	C	T	G	T	C	T	A	A	A	G	C	C	T	T	G	C	A		1801
Qy	3016	G	C	T	C	T	G	T	T	C	T	G	A	A	C	A	C	T	T	C	C	T	G	A	T	T	C	T	G	T	G	A	G	G	G	G	A	T	G	T	G	A	G	G	G		3075		
Db	1801	G	C	T	C	T	G	T	T	C	T	G	A	A	C	A	C	T	T	C	C	T	G	A	T	T	C	T	G	T	G	A	G	G	G	G	A	T	G	T	G	A	G	G	G		1860		
Qy	3076	A	A	G	A	G	G	C	A	G	C	T	G	A	G	C	C	A	C	A	G	G	G	A	G	G	T	G	A	G	G	G	G	A	C	A	G	A	A	G	G	C	A	G		3135			
Db	1861	A	A	G	A	G	G	C	A	G	C	T	G	A	G	C	C	A	C	A	G	G	G	A	G	T	G	A	G	G	G	G	A	C	A	G	A	A	G	G	C	A	G		1920				
Qy	3136	C	A	A	A	G	C	T	G	G	T	C	C	A	T	C	A	G	T	C	C	A	T	C	A	G	T	C	C	A	G	T	C	C	A	G	T	C	C	A	G	A	G		3195				
Db	1921	C	A	A	A	G	C	T	G	G	T	C	C	A	T	C	A	G	T	C	C	A	T	C	A	G	T	C	C	A	G	T	C	C	A	G	T	C	C	A	G	A	G		1980				
Qy	3196	C	A	A	T	C	T	T	C	A	G	A	A	G	C	T	C	A	A	T	T	C	C	T	C	C	T	C	C	T	C	C	T	C	C	T	C	C	T	C	A	A	G		3255				
Db	1981	C	A	A	T	C	T	T	C	A	G	A	A	G	C	T	C	A	A	T	T	C	C	T	C	C	T	C	C	T	C	C	T	C	C	T	C	C	T	C	A	A	G		2040				
Qy	3256	A	A	T	G	C	A	T	T	G	C	A	A	A	A	A	T	G	C	A	G	A	C	T	A	A	C	T	A	C	T	G	T	G	T	A	G	T	T	G	C	T	G		3315				
Db	2041	A	A	T	G	C	A	T	T	G	C	A	A	A	A	A	T	G	C	A	G	A	C	T	A	A	C	T	A	C	T	G	T	G	T	A	G	T	T	G	C	T	G		2100				
Qy	3316	G	C	A	T	T	C	A	A	A	A	A	A	A	A	A	T	G	C	A	G	A	T	T	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T		3375				
Db	2101	G	C	A	T	T	C	A	A	A	A	A	A	A	A	A	T	G	C	A	G	A	T	T	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T		2160					
Qy	3376	T	G	A	C	A	G	C	A	C	C	C	A	G	C	T	C	A	G	A	G	C	T	G	T	G	A	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C		3435				
Db	2161	T	G	A	C	A	G	C	A	C	C	C	A	G	C	T	C	A	G	A	G	C	T	G	T	G	A	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C		2220				
Qy	3436	A	G	G	G	A	G	A	A	A	A	A	A	A	A	A	G	A	G	A	G	A	G	T	A	T	A	G	C	A	A	A	A	A	A	A	A	A	A	A	A	A	A		3495				
Db	2221	A	G	G	G	A	G	A	A	A	A	A	A	A	A	A	G	A	G	A	G	A	G	T	A	T	A	G	C	A	A	A	A																

Qy	4095	CATTATTTCAAGTGGCTAAAGTTACTCTTCGACAGTTTGGGTATAATTAATTGGCTATTGC	4154
Db	2880	CATTATTTCAAGTGGCTAAAGTTACTCTTCGACAGTTTGGGTATAATTAATTGGCTATTGC	2939
Qy	4155	CATTTGCTTTTTGTTTTTTCTCCTTGCGTTTATAATGTAAACGACGGATTATTAACCTA	4214
Db	2840	CATTTGCTTTTTGTTTTTTCTCCTTGCGTTTATAATGTAAACGACGGATTATTAACCTA	2999
Qy	4215	CAGTCCAGAAGCCTGTGAATTTGAAATGAGAAAAAATAACAATTTTTGTTTTTACCACCT	4274
Db	3000	CAGTCCAGAAGCCTGTGAATTTGAAATGAGAAAAAATAACAATTTTTGTTTTTACCACCT	3059
Qy	4275	TCTTAACCTAAAATTAAACATTTTTATTCATTTCGGAATAGGCCATAAACTCAAAGTGGTAAT	4334
Db	3060	TCTTAACCTAAAATTAAACATTTTTATTCATTTCGGAATAGGCCATAAACTCAAAGTGGTAAT	3119
Qy	4335	AACAGTACCTGTGTGATTTTGTCAATACCAATAGAAAATCACAGACATTTTATCTATATTAC	4394
Db	3120	AACAGTACCTGTGTGATTTTGTCAATACCAATAGAAAATCACAGACATTTTATCTATATTAC	3179
Qy	4395	AGTTGTGCAGATACGTTGTGAAGTGAATAATTTATATCTOAAACTACTTTGAAATTAGAC	4454
Db	3180	AGTTGTGCAGATACGTTGTGAAGTGAATAATTTATATCTCAAACTACTTTGAAATTAGAC	3239
Qy	4455	CTCTGCTGCATCTTGTTTTTTAACATATTAATAAAACATGTTTAAAATTTTGCATATTTTG	4514
Db	3240	CTCTGCTGCATCTTGTTTTTTAACATATTAATAAAACATGTTTAAAATTTTGCATATTTTG	3299
Qy	4515	ATAATCATATTTCAATTATCATTTTGTTCCTTTGTAAATCTATATTTTATATATTGAAAAC	4574
Db	3300	ATAATCATATTTCAATTATCATTTTGTTCCTTTGTAAATCTATATTTTATATATTGAAAAC	3359
Qy	4575	ATCTTTCTGAGAAGTTCCCAGATTTTACCBAATGAGGTTCTTGGCATGCACACACA	4634
Db	3360	ATCTTTCTGAGAAGTTCCCAGATTTTACCBAATGAGGTTCTTGGCATGCACACACA	3419
Qy	4635	GAGTAAGAACTGATTTAGAGGCTTAAACATTTGAGCTGCCTGAGATGCAAGACTGAAAT	4694
Db	3420	GAGTAAGAACTGATTTAGAGGCTTAAACATTTGAGCTGCCTGAGATGCAAGACTGAAAT	3479
Qy	4695	TAGAAAGTTCTCCAAAGATACACAGTTGTTTTTAAAGCTAGGGGTGAGGGGGAAATCTG	4754
Db	3480	TAGAAAGTTCTCCAAAGATACACAGTTGTTTTTAAAGCTAGGGGTGAGGGGGAAATCTG	3539
Qy	4755	CCGCTTCTATAGGAATGCTCTCCCTGGAGCCTGGTAGGGTGCCTCTTGTGTTCTGGCT	4814
Db	3540	CCGCTTCTATAGGAATGCTCTCCCTGGAGCCTGGTAGGGTGCCTCTTGTGTTCTGGCT	3599
Qy	4815	GGCTGTATATTTTCTCTGCTCCCTGCTACGTCCTTAAAGGACTTGTTTGGATCTCCAAGTCC	4874
Db	3600	GGCTGTATATTTTCTCTGCTCCCTGCTACGTCCTTAAAGGACTTGTTTGGATCTCCAAGTCC	3659
Qy	4875	TAGCATAGTGCTCGGCACAGTCGAGGTTCTCAATGAGTTTGAGAGTGAATGAATATA	4934
Db	3660	TAGCATAGTGCTCGGCACAGTGAAGTCTCAATGAGTTTGAGAGTGAATGAATATA	3719
Qy	4935	AACTAGAAATATATCTCTGTGAAATCAGCACACAGTAGTCCCTGGTGTAAAGTGTGTATA	4994
Db	3720	AACTAGAAATATATCTCTGTGAAATCAGCACACAGTAGTCCCTGGTGTAAAGTGTGTATA	3779
Qy	4995	CGTGTGTGTGTGTGTGTGTGTGTGPAAAAACAGGTGGAGATATATAGGAACATAATT	5054
Db	3780	C--GTGTGTGTGTGTGTGTGTGTGPAAAAACAGGTGGAGATATATAGGAACATAATT	3837
Qy	5055	GGGGTATGGTGCATAAAAATTGGATGTCTCTTTTAAAAAGAAACTCCAAAACAGACTTCTG	5114
Db	3838	GGGGTATGGTGCATAAAAATTGGATGTCTCTTTTAAAAAGAAACTCCAAAACAGACTTCTG	3897
Qy	5115	GAAGGTATATTTTCTAAGAAATCTGTGGCAGCGTGAAGGC AACCCCCCTGTGCACAGCCC	5174
Db	3898	GAAGGTATATTTTCTAAGAAATCTGTGGCAGCGTGAAGGC AACCCCCCTGTGCACAGCCC	3957

[illegible]

RESULT 12

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US-10-278-698-808
; Sequence 808, Application US/10278698
; Publication No. US20050037344A1
; GENERAL INFORMATION:
;
; APPLICANT: PathoArray GmbH
; APPLICANT: Stuhlmüller, Bruno
; APPLICANT: Haupl, Thomas
; TITLE OF INVENTION: Nucleic Acid Array
; FILE REFERENCE: O30027US
; CURRENT APPLICATION NUMBER: US/10/278,698
; CURRENT FILING DATE: 2002-10-23
; NUMBER OF SEQ ID NOS: 1050
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 808
; LENGTH: 2800
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-278-698-808

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Query Match	34.2%	Score 1804.4	DB 8	Length 2800
Best Local Similarity	99.8%	Pred. No. 0		
Matches 1838	Conservative 0	Mismatches 1	Indels 3	Gaps 3
Qy	3431	AGCGCAGGGAGGAGAAAGAGAGAGGAGATAGTGTATGACGACAGAAAGACAGATTCA	T	3490
Db	1	AGCGCAGGGAGGAGAAAG-AAAAGAGAGGGATAGTGTATGACGAAAGAACAGATTCA	T	59
Qy	3491	TCAAGGCGAGTGGGAAATTGACCAACAGGATTTATAGTCCACGTGATCTCGTGGTCTAGGAG		3550
Db	60	TCAAGGCGAGTGGGAAATTGACCAACAGGATTTATAGTCCACGTGATCTCGTGGTCTAGGAG		119
Qy	3551	GCAGGGCTATATTGTGGGGGGGAAAAATCAGTTCAAGGGAAGTCGGGAGACCTGATTTCT		3610
Db	120	GCAGGGCTATATTGTGGGGGGGAAAAATCAGTTCAAGGGAAGTCGGGAGACCTGATTTCT		179
Qy	3611	AATACTATATTTTTTCCTTTCAAGCTGAGTAATTTCTGACGAGTCAAGGTTAGTAATCTG		3670
Db	180	AATACTATATTTTTTCCTTTCAAGCTGAGTAATTTCTGACGAGTCAAGGTTAGTAATCTG		239
Qy	3671	AGGCTGPAAGATTAATTAGTTTCTCCTTATAGGAATCTTTTTTCTCTGTGGAGTTAGCA		3730
Db	240	AGGCTGPAAGATTAATTAGTTTCTCCTTATAGGAATCTTTTTTCTCTGTGGAGTTAGCA		299


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Db 1319 ATCTGCCGCTCTAAGGAATGCTCTCCCTGGAGCTGTGAGGCTGTCTCTGTTGTC 1378
Qy 4810 TGGCTGGCTGTATTTTCTCTGCTCCCTGCTACGCTTTAAAGGACTTGTGTGGATCTCCA 4869
Db 1379 TGGCTGGCTGTATTTTCTCTGCTCCCTGCTACGCTTTAAAGGACTTGTGTGGATCTCCA 1438
Qy 4870 GTTCTAGCAGTAGTCCCTGGCACAGTGCAGGTTCTCAATGAGTTTGCAGAGTGAATGAA 4929
Db 1439 GTTCTAGCAGTAGTCCCTGGCACAGTGCAGGTTCTCAATGAGTTTGCAGAGTGAATGAA 1498
Qy 4930 ATATAAACTAGAAATATATCTTGTGTAATCAGCACACAGTGTCTCTGTTAAGTGT 4989
Db 1499 ATATAAACTAGAAATATATCTTGTGTAATCAGCACACAGTGTCTCTGTTAAGTGT 1558
Qy 4990 GTGTACGCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 5049
Db 1559 GTGTACGCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 1618
Qy 5050 TTATTGGGCTATGGGTGATATAAATGGGATGTTCTTTTAAAGGAACTCCAAACAGAC 5109
Db 1619 TTATTGGGCTATGGGTGATATAAATGGGATGTTCTTTTAAAGGAACTCCAAACAGAC 1678
Qy 5110 TTCTGGAGGTTATTTCTTAAGAACTTTGCTGGCAGCGTGAAGCAACCCCTGTGTGCAC 5169
Db 1679 TTCTGGAGGTTATTTCTTAAGAACTTTGCTGGCAGCGTGAAGCAACCCCTGTGTGCAC 1738
Qy 5170 AGCCCCACCCAGCTCAGTGGCAGCTCTGTCTTCCCTCCATGAAGGCTGGCTCCCGAG 5229
Db 1739 AGCCCCACCCAGCTCAGTGGCAGCTCTGTCTTCCCTCCATGAAGGCTGGCTCCCGAG 1798
Qy 5230 TATATATAAACCCTCTCTGGAGCTCGGGCATGAGCCAGG 5271
Db 1799 TATATATAAACCCTCTCTGGAGCTCGGGCATGAGCCAGG 1840

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RESULT 14

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US-10-956-243-1
; Sequence 1, Application US/10956243
; Publication No. US20050123960A1
; GENERAL INFORMATION:
; APPLICANT: Stone, Edwin M.
; APPLICANT: Sheffield, Val C.
; APPLICANT: Alward, Wallace L.M.
; APPLICANT: Fingert, John
; TITLE OF INVENTION: GLAUCOMA THERAPEUTICS AND DIAGNOSTICS
; FILE REFERENCE: 21087.0017U11
; CURRENT APPLICATION NUMBER: US/10/956,243
; CURRENT FILING DATE: 2004-10-01
; PRIOR APPLICATION NUMBER: US/09/952,464
; PRIOR FILING DATE: 2001-09-12
; PRIOR APPLICATION NUMBER: 09/473,273
; PRIOR FILING DATE: 1999-12-28
; PRIOR APPLICATION NUMBER: 09/461,542
; PRIOR FILING DATE: 1999-12-15
; PRIOR APPLICATION NUMBER: 09/366,952
; PRIOR FILING DATE: 1999-08-04
; PRIOR APPLICATION NUMBER: 09/056,285
; PRIOR FILING DATE: 1998-04-07
; PRIOR APPLICATION NUMBER: 08/822,999
; PRIOR FILING DATE: 1997-03-21
; NUMBER OF SEQ ID NOS: 43
; SOFTWARE: Fast-Seq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 2800
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence; Note =
; OTHER INFORMATION: Synthetic construct
US-10-956-243-1

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Query Match

34.2%; Score 1804.4; DB 9; Length 2800;

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Best Local Similarity 99.8%; Pred. No. 0;
Matches 1838; Conservative 0; Mismatches 1; Indels 3; Gaps 3;
Qy 3431 AGCCAGGGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 3490
Db 1 AGCCAGGGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 59
Qy 3491 TCAGGGCAGTGGGAATTTGACCCAGAGGATTTATAGTCCAGTGATCTCTGGGTTCTAGAG 3550
Db 60 TCAGGGCAGTGGGAATTTGACCCAGAGGATTTATAGTCCAGTGATCTCTGGGTTCTAGAG 119
Qy 3551 GCAGGGCTATATTGTGGGGGAGAAAAATCAGTTCAAGGAGAGTCCGGAGAGCTCATTTCT 3610
Db 120 GCAGGGCTATATTGTGGGGGAGAAAAATCAGTTCAAGGAGAGTCCGGAGAGCTCATTTCT 179
Qy 3611 AATACTATATTTTCTTTTCAAGCTCAGTAATTTCTGAGCAAGTCACAAGGTAGTAACCTG 3670
Db 180 AATACTATATTTTCTTTTCAAGCTCAGTAATTTCTGAGCAAGTCACAAGGTAGTAACCTG 239
Qy 3671 AGCTGTAAAGTACTTACTTCTCTTATAGGAGTCTTTTCTCTGTGGAGTTAGCA 3730
Db 240 AGCTGTAAAGTACTTACTTCTCTTATAGGAGTCTTTTCTCTGTGGAGTTAGCA 299
Qy 3731 GCACAGGGCAATCCCGTTTCTTTTAAACAGAGAGAAAAACATTCCTAAGAGTAAAGCCAAA 3790
Db 300 GCACAGGGCAATCCCGTTTCTTTTAAACAGAGAGAAAAACATTCCTAAGAGTAAAGCCAAA 359
Qy 3791 CAGATTCAAGCTTAGTCTTGTGACTATATGATTTGTTTTTGAAGAAATCATTTTCAGCG 3850
Db 360 CAGATTCAAGCTTAGTCTTGTGACTATATGATTTGTTTTTGAAGAAATCATTTTCAGCG 419
Qy 3851 ATGTTTACTATCTGATTCAGAAATGAGACTAGTACCTTTGTCAGCTGTAAACAACA 3910
Db 420 ATGTTTACTATCTGATTCAGAAATGAGACTAGTACCTTTGTCAGCTGTAAACAACA 479
Qy 3911 CCCAGTTGTAATGTCTCAAGTTTCAAGTTTCAAGTTTCAAGTTTCAAGTTTCAAGTTTCAAG 3969
Db 480 CCCAGTTGTAATGTCTCAAGTTTCAAGTTTCAAGTTTCAAGTTTCAAGTTTCAAGTTTCAAG 539
Qy 3970 TCTTTAGAGCAAACTGTGTTTCTCCACATCTCGAGGTGAGTCTGCCAGGCGAGTTTGGAA 4029
Db 540 TCTTTAGAGCAAACTGTGTTTCTCCAC-TCCTGAGGTGAGTCTGCCAGGCGAGTTTGGAA 598
Qy 4030 ATATTTACTTCAACAGTATGACATGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTG 4089
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Qy 4090 GCAATCATTTTCAAGTGGCTTAAAGTTTACTTCTGACAGTTTGGTATATTTTATTCGGCT 4149
Db 659 GCAATCATTTTCAAGTGGCTTAAAGTTTACTTCTGACAGTTTGGTATATTTTATTCGGCT 718
Qy 4150 ATTGCCATTTGTTTTTGTGTTTTTGTGTTTTTGTGTTTTTGTGTTTTTGTGTTTTTGTGTTTAC 4209
Db 719 ATTGCCATTTGTTTTTGTGTTTTTGTGTTTTTGTGTTTTTGTGTTTTTGTGTTTTTGTGTTTAC 778
Qy 4210 ACCACAGTCCAGAAAGCCTGTAATTTGAAATGAGGAGAAAAATATATTTTGTGTTTTTAC 4269
Db 779 ACCACAGTCCAGAAAGCCTGTAATTTGAAATGAGGAGAAAAATATATTTTGTGTTTTTAC 838
Qy 4270 CACCTTTCTAAATTTAAACATTTTATTTCCATTTGGAATAGAGCCATAAATCTCAAGTG 4329
Db 839 CACCTTTCTAAATTTAAACATTTTATTTCCATTTGGAATAGAGCCATAAATCTCAAGTG 898
Qy 4330 GTAATAACAGTACCTGTGATTTTGTGTTTATTAATGTAAGAGGAGTTATTA 4389
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Db 959 ATTACAGTTGTTGAGATACCTGTTGAAGTGAATTTTATCTCAAACTACTTCAAAAT 1018
Qy 4450 TAGACCTCTGCTGGATCTTTGTTTTTAAACATTTTAAACATTTTAAATTTTGTATA 4509

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Db 1019 TAGACCTCTGCTGGAGTCTTTTAAACATATTAATAAACAATGTTTAAATTTGATA 1078
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Qy 4570 AAAACATCTTTCTGAGAGAGTCTCCAGATTTTCCCAATGAGGTTCTTGGCATGCAC 4629
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Qy 4690 GAAATAGAAAGTTCTCCCAAGATACACAGTTGTTTAAAGCTAGAGGCTGAGGGGAA 4749
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Qy 4750 ATCTGCGCTTCTATAGGAATGCTCTCCCTGGAGCCTGGTAGGGTGTCTTGTGTTTC 4809
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Qy 4810 TGGCTGGCTGTTATTTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4869
Db 1379 TGGCTGGCTGTTATTTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1438
Qy 4870 GTTCTAGCATAGTGGCTGGCAGAGTCTCAATGAGTTTGCAGAGTGAATGGAA 4929
Db 1439 GTTCTAGCATAGTGGCTGGCAGAGTCTCAATGAGTTTGCAGAGTGAATGGAA 1498
Qy 4930 ATATAAAGTAAATATATCTTGTGTAATTCAGACACACAGTAGTCTGCTGCTGCTGCT 4989
Db 1499 ATATAAAGTAAATATATCTTGTGTAATTCAGACACACAGTAGTCTGCTGCTGCTGCT 1558
Qy 4990 GTCTAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5049
Db 1559 GTCTAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1618
Qy 5050 TTATTTGGGTTATGGGTCATAAATTTGGGATGTTCTTTTAAAGAACTCCAAACAGAC 5109
Db 1619 TTATTTGGGTTATGGGTCATAAATTTGGGATGTTCTTTTAAAGAACTCCAAACAGAC 1678
Qy 5110 TTCTGGAAGTTATTTTCTAAGAACTTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5169
Db 1679 TTCTGGAAGTTATTTTCTAAGAACTTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1738
Qy 5170 AGCCCCCAGCCTCAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5229
Db 1739 AGCCCCCAGCCTCAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1798
Qy 5230 TATATATAAACCCTCTCTGGAGCTGGGCTGAGCCAGCAAGG 5271
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RESULT 15

US-09-925-065A-905501
; Sequence 905501, Application US/09925065A
; Publication No. US20050228172A9
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single
; TITLE OF INVENTION: Nucleotide Polymorphisms in the Human Genome
; FILE REFERENCE: 108827.135
; CURRENT APPLICATION NUMBER: US/09/925,065A
; CURRENT FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: US 60/243,096
; PRIOR FILING DATE: 2000-10-24
; PRIOR APPLICATION NUMBER: US 60/252,147
; PRIOR FILING DATE: 2000-11-20
; PRIOR APPLICATION NUMBER: US 60/250,092
; PRIOR FILING DATE: 2000-11-30
; PRIOR APPLICATION NUMBER: US 60/261,766

; PRIOR FILING DATE: 2001-01-16
; PRIOR APPLICATION NUMBER: US 60/289,846
; PRIOR FILING DATE: 2001-05-09
; NUMBER OF SEQ ID NOS: 957086
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 905501
; LENGTH: 632
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-925-065A-905501

Query Match 12.0%; Score 631.6; DB 4; Length 632;
Best Local Similarity 99.8%; Pred. No. 1.1e-145;
Matches 631; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1284 TGAGGCTCTGCTCTTACACCTACCTGATGCTTACACCTGAGTCACTGCAACCTCTG 1343
Db 1 TGAGGCTCTGCTCTTACACCTACCTGATGCTTACACCTGAGTCACTGCAACCTCTG 60
Qy 1344 CCTCCAGGTTCAAGCAATTCCTGCTCAGCCTCCCGGTAGCTGGGACTACAGGCG 1403
Db 61 CCTCCAGGTTCAAGCAATTCCTGCTCAGCCTCCCGGTAGCTGGGACTACAGGCG 120
Qy 1404 AGCCCGGCTAAATTTTGTATTGTAGTAGAGATGGGTTTCAACATATTAGCCCGCTG 1463
Db 121 AGCCCGGCTAAATTTTGTATTGTAGTAGAGATGGGTTTCAACATATTAGCCCGCTG 180
Qy 1464 GTCTTGAATCTCTGACCTCAGGTGATCCACCCACCTCAGCCTCTTAAAGTGGGATTA 1523
Db 181 GTCTTGAATCTCTGACCTCAGGTGATCCACCCACCTCAGCCTCTTAAAGTGGGATTA 240
Qy 1524 CAGGATGATGATCAGCGCGCCCGGCTAGGTTTAAAGGAACTTGAATGG 1583
Db 241 CAGGATGATGATCAGCGCGCCCGGCTAGGTTTAAAGGAACTTGAATGG 300
Qy 1584 TTTTACTAAACCAAGGAAACAGACAAAGCTGTGATAATTCAGGGATTTCTGGGATG 1643
Db 301 TTTTACTAAACCAAGGAAACAGACAAAGCTGTGATAATTCAGGGATTTCTGGGATG 360
Qy 1644 GGGAAATGGTGCATGAGTGCCTGCTAGTCCAGACCACTGCTCTCATCTTCTTC 1703
Db 361 GGGAAATGGTGCATGAGTGCCTGCTAGTCCAGACCACTGCTCTCATCTTCTTC 420
Qy 1704 CCTCATCTCTCATTTTCAGGCTAAGTTACCATTTTATTCACCATGCTTTTGTGTAAGCCT 1763
Db 421 CCTCATCTCTCATTTTCAGGCTAAGTTACCATTTTATTCACCATGCTTTTGTGTAAGCCT 480
Qy 1764 CCACATCTGTTACTGAAATAAGAGTATACATAAACTAGTTTCCATTTGGGGCCATCTGTG 1823
Db 481 CCACATCTGTTACTGAAATAAGAGTATACATAAACTAGTTTCCATTTGGGGCCATCTGTG 540
Qy 1824 TGTGTATAGGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1883
Db 541 TGTGTATAGGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 600
Qy 1884 TCTCCAGCTGGGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 1915
Db 601 TCTCCAGCTGGGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 632

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Job time : 2632 secs

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: January 26, 2006, 04:13:16 ; Search time 514 Seconds
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Title: US-09-227-881-34

Perfect score: 5271

Sequence: 1 atcttgttcagttacctc.....tcgggcacgagccagcaagg 5271

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext.1.0

Searched: 6059551 seqs, 415333918 residues

Total number of hits satisfying chosen parameters: 12119102

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications NA New:

- 1: /cgm2_6/ptodata/1/pubpna/us08 NEW PUB.seq.*
- 2: /cgm2_6/ptodata/1/pubpna/us06 NEW PUB.seq.*
- 3: /cgm2_6/ptodata/1/pubpna/us07 NEW PUB.seq.*
- 4: /cgm2_6/ptodata/1/pubpna/PCT_NEW PUB.seq.*
- 5: /cgm2_6/ptodata/1/pubpna/us09 NEW PUB.seq.*
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- 11: /cgm2_6/ptodata/1/pubpna/us11 NEW PUB.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	190.8	3.6	16525	US-11-136-527-717	Sequence 717, App
2	185.4	3.5	86361	US-10-995-561-13364	Sequence 13364, A
3	185.2	3.5	143389	US-11-112-908-30	Sequence 30, Appl
4	185.2	3.5	150314	US-11-112-908-24	Sequence 24, Appl
5	185.2	3.5	166020	US-11-112-908-28	Sequence 28, Appl
6	184	3.5	60754	US-10-995-561-13440	Sequence 13440, A
7	183.4	3.5	415117	US-10-995-561-13274	Sequence 13274, A
8	179.6	3.4	14804	US-10-995-561-13379	Sequence 13379, A
9	179.6	3.4	98716	US-10-995-561-13331	Sequence 13331, A
10	179.6	3.4	191091	US-11-121-086-60	Sequence 60, Appl
11	179	3.4	159497	US-11-112-908-61	Sequence 61, Appl
12	179	3.4	171427	US-11-112-908-60	Sequence 60, Appl
13	178.4	3.4	127917	US-10-775-169-82	Sequence 82, Appl
14	177.8	3.4	161994	US-11-112-908-57	Sequence 57, Appl
15	177.8	3.4	168656	US-11-112-908-59	Sequence 59, Appl
16	177.8	3.4	170285	US-11-112-908-58	Sequence 58, Appl
17	177.8	3.4	387780	US-10-995-561-13259	Sequence 13259, A
18	177.6	3.4	128978	US-10-775-169-345	Sequence 345, App
19	176.8	3.4	53332	US-10-786-085-3	Sequence 3, Appl
20	176.8	3.4	179777	US-11-121-086-106	Sequence 106, App
21	176.6	3.4	1154	US-11-145-703-37	Sequence 37, Appl
22	176.6	3.4	153142	US-11-121-086-27	Sequence 27, Appl

c 23	176.2	3.3	1301	8	US-11-145-703-36	Sequence 36, Appl
c 24	176.2	3.3	1386	8	US-11-145-703-40	Sequence 40, Appl
c 25	176	3.3	207600	8	US-11-112-908-31	Sequence 31, Appl
c 26	176	3.3	139608	8	US-11-145-703-1	Sequence 1, Appl
c 27	175.6	3.3	177623	8	US-11-112-908-41	Sequence 41, Appl
c 28	175	3.3	86081	7	US-10-995-561-13246	Sequence 13246, A
c 29	174.8	3.3	17455	7	US-10-995-561-13426	Sequence 13426, A
c 30	174.8	3.3	28724	7	US-10-995-561-13372	Sequence 13372, A
c 31	174.8	3.3	155989	8	US-11-121-086-57	Sequence 57, Appl
c 32	174.8	3.3	180654	8	US-11-121-086-58	Sequence 58, Appl
c 33	174.6	3.3	40987	7	US-10-995-561-13503	Sequence 13503, A
c 34	174.6	3.3	67088	7	US-10-995-561-13365	Sequence 13365, A
c 35	174.6	3.3	79528	7	US-10-276-233A-6	Sequence 6, Appl
c 36	174.4	3.3	60844	7	US-10-995-561-13359	Sequence 13359, A
c 37	174.2	3.3	115935	7	US-10-775-169-241	Sequence 241, App
c 38	174.2	3.3	141121	7	US-10-995-561-13262	Sequence 13262, A
c 39	174.2	3.3	162289	8	US-11-121-086-20	Sequence 20, Appl
c 40	174.2	3.3	191684	8	US-11-121-086-2	Sequence 2, Appl
c 41	174.2	3.3	197096	8	US-11-121-086-107	Sequence 107, App
c 42	174	3.3	13609	7	US-10-995-561-13442	Sequence 13442, A
c 43	174	3.3	16082	7	US-10-995-561-13485	Sequence 13485, A
c 44	174	3.3	23894	7	US-10-995-561-13320	Sequence 13320, A
c 45	174	3.3	87672	7	US-10-995-561-13237	Sequence 13237, A

ALIGNMENTS

RESULT 1

US-11-136-527-717
; Sequence 717, Application US/11136527
; Publication No. US20050287570A1
; GENERAL INFORMATION:
; APPLICANT: Wyeth
; APPLICANT: Mounts, William M
; TITLE OF INVENTION: Probe Arrays For Expression Profiling of Rat Genes
; FILE REFERENCE: 031896-041000 (AM101086)
; CURRENT APPLICATION NUMBER: US/11/136.527
; CURRENT FILING DATE: 2005-05-25
; PRIOR APPLICATION NUMBER: US 60/574,294
; PRIOR FILING DATE: 2005-05-26
; NUMBER OF SEQ ID NOS: 362830
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 717
; LENGTH: 16525
; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-11-136-527-717

Query Match	3.6%	Score 190.8;	DB 8;	Length 16525;
Best Local Similarity	55.9%	Pred No. 6.1e-33;		
Matches 871;	Conservative 0;	Mismatches 542;	Indels 145;	Gaps 21;
Qy	2387	CTTGGCTCCAGGCTCCAGAAAGGAAATGGAGAGGAAACTAGTCTAAACGGAGAATCTGGA	2446	
Db	2815	CCTGGCTCCTGGCTCCAGAAAGGATATGGAGATAGTCTCCTCGTAACATTGGA	2874	
Qy	2447	GGGACAGTCTTCTCCTCAGAGGAAAGGGCCCTCCAGTCCAGGAGATTCAGAGGTG	2506	
Db	2875	ATGCCACAGTCTTCTCCTCAGTGGGAAAGAGTCTCTGTGTC	2931	
Qy	2507	GGGACTCAGGAGTGGGACGCTGGGGCTGCTGAAAGGCAGGAGGTGAAA	2566	
Db	2932	GGGACTCAGGAGGAGGCTGCTG	2981	
Qy	2567	AGGCAAGGCTGAAGCTGCCAGATGTTTCAGTGTGTTTCACGGGGCTGGGAGTTTCGGT	2626	
Db	2982	AGGG-TAGGCTAAATCTCACCAGATGTTTCGCTGCTGCTGGAGTAGGAGGCTCCTGT	3040	
Qy	2627	TGCTTCTGTGAGGCTTTTATCTTTCTCTGCTTGAGGAGAGAGTCTATTTCATGA	2686	
Db	3041	TTCTCCACAGGGCC-----CATTTCTGGGCTTAGAGGAGAACGTTTCTATTCTATGA	3093	

PRIOR FILING DATE: 2004-04-23
PRIOR APPLICATION NUMBER: US 60/575,978
PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US 60/631,702
PRIOR FILING DATE: 2004-11-30
PRIOR APPLICATION NUMBER: US 60/633,826
PRIOR FILING DATE: 2004-12-07
NUMBER OF SEQ ID NOS: 511
SOFTWARE: PatentIn version 3.3
SEQ ID NO 30
LENGTH: 143389
TYPE: DNA
ORGANISM: Homo sapiens
US-11-112-908-30

Query Match 3.5%; Score 185.2; DB 8; Length 143389;
Best Local Similarity 79.4%; Pred. No. 2.7e-31;
Matches 235; Conservative 0; Mismatches 53; Indels 8; Gaps 1;

QY 1280 AGGGTGAGGCTGTGTCTTACACCTACCTGATGCTCTACACCTGAGCTCACTGCAACC 1339
DB 113274 AGAGTGTGCTGTGTACCCAGATTGGAGTGGAGTGGACATCTCAGCTCACCGCAACC 113333

QY 1340 TCTGCTCCAGGTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGGACTACAG 1399
DB 113334 TCCGCTCCAGGTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGGACTACAG 113393

QY 1400 GCG-----CACGCCCGGCTAAATTTTGTATTGTATTGTAGTAGAGATGGGTTTCAACATA 1451
DB 113394 GCGCCCGCCACACCGCTGGCTAAATTTTGTATTGTATTGTAGTAGAGATGGGTTTCAACATA 113453

QY 1452 TTAGCCCGGCTGTGTCTTGAACCTTCCAGCTCAGGTGATCCACCCACCTCAGCTCTCTAAA 1511
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QY 1512 GTGCTGGGATTACAGGATGATGATCAGCGCCCGGCAAGGTCAGTGTATTATAA 1567
DB 113514 GTGCTGGGATTACAGGATGATGATCAGCGCCCGGCAAGGTCAGTGTATTATAA 113569

RESULT 4
US-11-112-908-24
Sequence 24, Application US/11112908
Publication No. US2005026059A1
GENERAL INFORMATION:
APPLICANT: Harris, Cole
TITLE OF INVENTION: Breast Cancer Biomarkers
FILE REFERENCE: 04-164-US
CURRENT APPLICATION NUMBER: US/11/112,908
PRIOR FILING DATE: 2005-04-22
PRIOR APPLICATION NUMBER: US 60/564,758
PRIOR FILING DATE: 2004-04-23
PRIOR APPLICATION NUMBER: US 60/631,702
PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US 60/575,978
PRIOR FILING DATE: 2004-11-30
PRIOR APPLICATION NUMBER: US 60/633,826
PRIOR FILING DATE: 2004-12-07
NUMBER OF SEQ ID NOS: 511
SOFTWARE: PatentIn version 3.3
SEQ ID NO 24
LENGTH: 150314
TYPE: DNA
ORGANISM: Homo sapiens
US-11-112-908-24

Query Match 3.5%; Score 185.2; DB 8; Length 150314;
Best Local Similarity 79.4%; Pred. No. 2.8e-31;
Matches 235; Conservative 0; Mismatches 53; Indels 8; Gaps 1;

QY 1280 AGGGTGAGGCTGTGTCTTACACCTACCTGATGCTCTACACCTGAGCTCACTGCAACC 1339

DB 27741 AGAGTGTGCTGTGTACCCAGATTGGAGTGGAGTGGACATCTCAGCTCACCGCAACC 27800

QY 1340 TCTGCTCCAGGTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGGACTACAG 1399
DB 27801 TCCGCTCCAGGTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGGACTACAG 27860

QY 1400 GCG-----CACGCCCGGCTAAATTTTGTATTGTATTGTAGTAGAGATGGGTTTCAACATA 1451
DB 27861 GCGCCCGCCACACCGCTGGCTAAATTTTGTATTGTATTGTAGTAGAGATGGGTTTCAACATA 27920

QY 1452 TTAGCCCGGCTGTGTCTTGAACCTTCCAGCTCAGGTGATCCACCCACCTCAGCTCTCTAAA 1511
DB 27921 TTGGTCAGGCTGTGTGTGACTCTTGAACCTTCCAGCTCAGGTGATCCACCCACCTCAGCTCTCTAAA 27980

QY 1512 GTGCTGGGATTACAGGATGATGATCAGCGCCCGGCAAGGTCAGTGTATTATAA 1567
DB 27981 GTGCTGGGATTACAGGATGATGATCAGCGCCCGGCAAGGTCAGTGTATTATAA 28036

RESULT 5
US-11-112-908-28
Sequence 28, Application US/11112908
Publication No. US2005026059A1
GENERAL INFORMATION:
APPLICANT: Harris, Cole
TITLE OF INVENTION: Breast Cancer Biomarkers
FILE REFERENCE: 04-164-US
CURRENT APPLICATION NUMBER: US/11/112,908
PRIOR FILING DATE: 2005-04-22
PRIOR APPLICATION NUMBER: US 60/564,758
PRIOR FILING DATE: 2004-04-23
PRIOR APPLICATION NUMBER: US 60/575,978
PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US 60/631,702
PRIOR FILING DATE: 2004-11-30
PRIOR APPLICATION NUMBER: US 60/633,826
PRIOR FILING DATE: 2004-12-07
NUMBER OF SEQ ID NOS: 511
SOFTWARE: PatentIn version 3.3
SEQ ID NO 28
LENGTH: 166020
TYPE: DNA
ORGANISM: Homo sapiens
US-11-112-908-28

Query Match 3.5%; Score 185.2; DB 8; Length 166020;
Best Local Similarity 79.4%; Pred. No. 2.9e-31;
Matches 235; Conservative 0; Mismatches 53; Indels 8; Gaps 1;

QY 1280 AGGGTGAGGCTGTGTCTTACACCTACCTGATGCTCTACACCTGAGCTCACTGCAACC 1339
DB 159755 AGAGTGTGCTGTGTACCCAGATTGGAGTGGAGTGGACATCTCAGCTCACCGCAACC 159814

QY 1340 TCTGCTCCAGGTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGGACTACAG 1399
DB 159815 TCCGCTCCAGGTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGGACTACAG 159874

QY 1400 GCG-----CACGCCCGGCTAAATTTTGTATTGTATTGTAGTAGAGATGGGTTTCAACATA 1451
DB 159875 GCGCCCGCCACACCGCTGGCTAAATTTTGTATTGTATTGTAGTAGAGATGGGTTTCAACATA 159934

QY 1452 TTAGCCCGGCTGTGTCTTGAACCTTCCAGCTCAGGTGATCCACCCACCTCAGCTCTCTAAA 1511
DB 159935 TTGGTCAGGCTGTGTGTGACTCTTGAACCTTCCAGCTCAGGTGATCCACCCACCTCAGCTCTCTAAA 159994

QY 1512 GTGCTGGGATTACAGGATGATGATCAGCGCCCGGCAAGGTCAGTGTATTATAA 1567
DB 159995 GTGCTGGGATTACAGGATGATGATCAGCGCCCGGCAAGGTCAGTGTATTATAA 160050

RESULT 6
US-10-995-561-13440/C

Sequence 13440, Application US/10995561
Publication No. US2005027054A1
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 13440
LENGTH: 60754
TYPE: DNA
ORGANISM: Homo sapiens
US-10-995-561-13440

Query Match 3.5%; Score 184; DB 7; Length 60754;
Best Local Similarity 71.9%; Pred. No. 3.6e-31;
Matches 256; Conservative 1; Mismatches 91; Indels 8; Gaps 1;

QY 1283 GTGAGGCTGTCTTACACCTACCTGTATGCTCTACACCTGAGCTCACTGCAACCTCT 1342
DB 9703 GTCTCGCTGTGCGCCAGGCTGGCATGAGTGAGTCTCAGCTCACTGCAACCTCT 9644

QY 1343 GCCTCCAGGTTCAAGCAATCTCTCTCAGCCTCCCGCTGAGTGGGACTACAGCG 1402
DB 9643 GTCTCCCGGGTTCAAGGATTTTCTGCTCAGCTCCAGTGTAGTGGGACTACAGGAG 9584

QY 1403 C-----ACGCCGGCTAAATTTTGTATTGTAGTGGGTTTCAACATTA 1454
DB 9583 CTGTCACTATGCTGGCTAAATTTTATATTTTAGTAGAGATGGGGTTTCAAGATG 9524

QY 1455 GCCCGGCTGTCTTGAACCTCTCAGCTCAGTGTATCCAGCTCAGCTCCTCTAAATG 1514
DB 9523 GCCAGGCTGTCTTGAACCTCTCAGCTCAGTGTATCCAGCTCAGCTCCTCTAAATG 9464

QY 1515 CTGGGATTACAGGATGAGTACCGCGCCCGGCAAGGTCAGTGTATTAAGGAATA 1574
DB 9463 CTGGGATTACAGGATGAGTACCGCGCCCGGCAAGTCAATTAACAAACAA 9404

QY 1575 CTTGATGTTTACTTAACCAACAGGGAACACACAAAGCTGTATATTTTCAGG 1630
DB 9403 TAATAATAATAATAAAGAAAGAAACAAACAAATATACAGATATATACGGG 9348

RESULT 7
US-10-995-561-13274
Sequence 13274, Application US/10995561
Publication No. US2005027054A1
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 13274
LENGTH: 41517
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(41517)
OTHER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables 1-
US-10-995-561-13274

Query Match 3.5%; Score 183.4; DB 7; Length 41517;
Best Local Similarity 76.5%; Pred. No. 1.1e-30;

Matches 241; Conservative 0; Mismatches 66; Indels 8; Gaps 1;

QY 1280 AGGGTGGAGGCTGTGTCTTACACCTACCTGTATGCTCTACACCTGAGCTCACTGCAAC 1339
DB 60473 AGAGTCTTGTCTGTCAACCCAGCTGTAGTGTCAATGGCGCAATCTCAGCTCACTGCAAC 60532

QY 1340 TCTGCTCCAGGTTCAAGCAATCTCTCTCAGCCTCCCGCTGAGTGGGACTACAG 1399
DB 60533 TCCGCTCCAGGTTCAAGTGTATCTCTGCTCAGCTCTCTGAGTAGTGGGATACAG 60592

QY 1400 GCG-----CACGCCCGGCTAAATTTTGTATTGTATGTAGTAGAGTGGGTTTCAACATA 1451
DB 60593 GTGAGTGGCACCACCTCCCGGCTAAATTTTGTATTGTATGTAGTAGAGTGGGTTTACCATG 60652

QY 1452 TTAGCCCGGCTGTGTGAACTCTGAGCTCAGGTGATCCACCCACCTCAGCTCCTTAA 1511
DB 60653 TTGGCCAGGCTGTGTGAACTCTGAGCTCAGGTGATCCACCCACCTTGGGCTTCCCAA 60712

QY 1512 GTGCTGGGATTACAGGCTAGTACCGCGCCCGGCTGAGTGTATTAAGGA 1571
DB 60713 GTGCTGGGATTACAGGCTAGTACCGCGCCCGGCTGAGTGTATTAAGGA 60772

QY 1572 TAACTTGAATGTTT 1586
DB 60773 TGTCTGGAGAGCTT 60787

RESULT 8
US-10-995-561-13379
Sequence 13379, Application US/10995561
Publication No. US2005027054A1
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 13379
LENGTH: 14804
TYPE: DNA
ORGANISM: Homo sapiens
US-10-995-561-13379

Query Match 3.4%; Score 179.6; DB 7; Length 14804;
Best Local Similarity 80.0%; Pred. No. 1.9e-30;
Matches 224; Conservative 2; Mismatches 46; Indels 8; Gaps 1;

QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGCTCAGGCTCC 1380
DB 13508 ATCTCAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGCTCAGGCTCC 13567

QY 1381 CGCGTAGCTGGGACTACAGGCT-----GCAGCCCGGCTAAATTTTGTATTGTAGTA 1432
DB 13568 CAAGTAGCTGGGATTACAGGCTACAGGCTACAGGCTACAGGCTACAGGCTACAGGCT 13627

QY 1433 GAGATGGGTTTCAACCATATTAGCCCGGCTGTGTTGAACCTCTGACCTCAGGTATCCA 1492
DB 13628 GAGACGGGTTTCAACCATATTAGCCCGGCTGTGTTGAACCTCTGACCTCAGGTATCCA 13687

QY 1493 CCCACCTCAGCTCTTAAAGTGTGGGATTACAGGCTACAGGCTACAGGCTACAGGCTACAGG 1552
DB 13688 CCCACCTCAGCTCTTAAAGTGTGGGATTACAGGCTACAGGCTACAGGCTACAGGCTACAGG 13747

QY 1553 GTCAGTGTATTAAGGATTAAGTGTGTTTACTTAA 1592
DB 13748 TATTTCAKTTAAGCTCAATAATCAATAATGAGTAGCA 13787

RESULT 9

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US-10-995-561-13331/c
; Sequence 13331, Application US/10995561
; Publication No. US20050272054A1
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
; TITLE OF INVENTION: DETECTION AND USES THEREOF
; FILE REFERENCE: CL001559
; CURRENT APPLICATION NUMBER: US/10/995,561
; CURRENT FILING DATE: 2004-11-24
; NUMBER OF SEQ ID NOS: 85702
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13331
; LENGTH: 98716
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-995-561-13331

Query Match 3.4%; Score 179.6; DB 7; Length 98716;
Best Local Similarity 80.0%; Pred. No. 4.3e-30;
Matches 224; Conservative 2; Mismatches 46; Indels 8; Gaps 1;

Qy 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTCCTCTGCTCAGCCTCC 1380
Db 75944 ATCTCAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTCCTCTGCTCAGCCTCC 75885

Qy 1381 CGCTAGCTGGGACTACAGGC-----GCAGCCCGGCTAAATTTTGTATTTAGTA 1432
Db 75884 CAAGTAGCTGGGATTACAGGCAGCACCACAGCCAGTAAATTTTGTATTTCTAGTA 75825

Qy 1433 GAGATGGGGTTTACCATTATAGCCGCTGCTTGAATCTGACCTCAGGTGATCCA 1492
Db 75824 GAGACGGGGTTTACCATTATAGCCGCTGCTTGAATCTGACCTCAGGTGATCCA 75765

Qy 1493 CCCACCTCAGCTCTCTAAAGTCTGGGATTACAGGCATGAGTCACCGCGCCCGGCAAGG 1552
Db 75764 CCCACCTCGGTCTCCAAAGTCTGGGATTACAGGCATGAGTCACCGCGCCCGGCGT 75705

Qy 1553 GTCAGTGTATTAAGGAATACCTGAATGTTTACTAAA 1592
Db 75704 TATTTCAKTTAAGCTCAAAATACAAATATGAGTAAGCAA 75665

RESULT 10
US-11-121-086-60
; Sequence 60, Application US/11121086
; Publication No. US20050266459A1
; GENERAL INFORMATION:
; APPLICANT: POULSEN, TIM S.
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES
; FILE REFERENCE: 09138.6000-0000
; CURRENT APPLICATION NUMBER: US/11/121,086
; CURRENT FILING DATE: 2005-05-04
; PRIOR APPLICATION NUMBER: 60/567,570
; PRIOR FILING DATE: 2004-05-04
; NUMBER OF SEQ ID NOS: 107
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 60
; LENGTH: 191091
; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-121-086-60

Query Match 3.4%; Score 179.6; DB 8; Length 191091;
Best Local Similarity 79.9%; Pred. No. 5.6e-30;
Matches 227; Conservative 0; Mismatches 49; Indels 8; Gaps 1;

Qy 1290 TCTGTGTTTACACTACCTGATGCTTACCTGAGCTCACTGCAACCTCTGCTCCCTCC 1349
Db 104701 TCTGTCCCTAGGCTGGAGTGCAGTGGGCAAAATCTTAGCTCACTGCAACCTCTCC 104760

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Qy 1350 AGTTTCAAGCAATTCCTCTGCTCAGCCTCCGCGTAGCTGGGACTACAGGC----- 1401
Db 104761 AGTTTCAAGCAATTCCTCTGCTCAGCCTCCGCGTAGCTGGGACTACAGGCACACACTA 104820

Qy 1402 GCAGCCCGGCTAAATTTTGTATTTAGTAGAGATGGGGTTTACACCATATTAGCCCGC 1461
Db 104821 TCAGCCCATCTAAATTTTGTATTTTGTAGTGAGACAGGGTTTACACCATATTAGCAG 104880

Qy 1462 TGGTCTTGAACCTCTGACCTCAGGTGATCCACCCACCTCAGCCTCTTAAAGTGTGGGAT 1521
Db 104881 TGGTCTCGAATCTCTGACCTCAGGTGATCCACCTCTCGGCTCCCAAGTGTGGGAT 104940

Qy 1522 TACAGGCATGAGTCACCGCGCCGCGCCCAAGGTCTAGTGTATTAAT 1565
Db 104941 TACAGGCATGAGTCACCGCGCCGCGCCCAATGTCTAGTGTATTAAT 104984

RESULT 11
US-11-112-908-61/c
; Sequence 61, Application US/11112908
; Publication No. US20050260659A1
; GENERAL INFORMATION:
; APPLICANT: Harris, Lisa M.
; TITLE OF INVENTION: Breast Cancer Biomarkers
; FILE REFERENCE: 04-164-US
; CURRENT APPLICATION NUMBER: US/11/112,908
; CURRENT FILING DATE: 2005-04-22
; PRIOR APPLICATION NUMBER: US 60/564,758
; PRIOR FILING DATE: 2004-04-23
; PRIOR APPLICATION NUMBER: US 60/575,978
; PRIOR FILING DATE: 2004-06-01
; PRIOR APPLICATION NUMBER: US 60/631,702
; PRIOR FILING DATE: 2004-11-30
; PRIOR APPLICATION NUMBER: US 60/633,826
; PRIOR FILING DATE: 2004-12-07
; NUMBER OF SEQ ID NOS: 511
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 61
; LENGTH: 159497
; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-112-908-61

Query Match 3.4%; Score 179; DB 8; Length 159497;
Best Local Similarity 81.0%; Pred. No. 7.1e-30;
Matches 221; Conservative 0; Mismatches 50; Indels 2; Gaps 1;

Qy 1281 GGGTGAGGCTCTGTGCTTACACCTACCTGATGCTTACACCTGAGCTCACTGCAACCT 1340
Db 100954 GAGTGTGCTCTGTCAACTAGGCTGGGATGAGTGGGATGATCTCAGCTCACTGCAACCT 100895

Qy 1341 CTGCTCCCAAGGTTCAAGCAATTCCTCTGCTCAGCCTCCGCGTAGCTGGGACTACAGG 1400
Db 100894 CTGCTCTCTGGTTCAAGCATTCCTCTGCTCAGCCTCCCAAGTAGTGGGATTACAGG 100835

Qy 1401 C--GCAGCCCGGCTAAATTTTGTATTTGTAGTAGAGATGGGGTTTACCATATTAGCCC 1458
Db 100834 CCACCATGCGCGCTAAATTTTGTATTTTGTAGTAGAGATGGGGTTTACCATATTAGGCA 100775

Qy 1459 GGCTGGTCTTGAACCTCTGACCTCAGGTGATCCACCCACCTCAGCCTCTTAAAGTGTGG 1518
Db 100774 GGCTGGTCTGAACTCTCTGACCTCAAGTATCACTCGCTCGGCTCCCAAGTGTGG 100715

Qy 1519 GATTACAGGCATGAGTCACCGCGCCCGCCCAAG 1551
Db 100714 GATTACAGGCATGAGTCACCGCGCTCGGCTCGGCTATG 100682

RESULT 12
US-11-112-908-60/c
; Sequence 60, Application US/11112908
; Publication No. US20050260659A1

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GenCore version 5.1.6
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SUMMARIES

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3	5271	100.0	6169	19	US-08-938-669-3	Sequence 3, Appl
4	5271	100.0	6169	22	US-09-227-881-3	Sequence 3, Appl
5	5271	100.0	6169	43	US-10-244-633-3	Sequence 3, Appl
6	5246.4	99.5	5300	19	US-08-938-669-1	Sequence 1, Appl
7	5246.4	99.5	5300	22	US-09-227-881-1	Sequence 1, Appl
8	5246.4	99.5	5300	38	US-09-985-673A-1	Sequence 1, Appl
9	5246.4	99.5	5300	43	US-10-244-633-1	Sequence 1, Appl
10	5246.4	99.5	5300	61	US-10-741-339-1	Sequence 1, Appl
11	5232.4	99.3	37252	40	US-10-087-192-1228	Sequence 1228, Ap
12	5224.4	99.1	5304	19	US-08-938-669-2	Sequence 2, Appl
13	5224.4	99.1	5304	22	US-09-227-881-2	Sequence 2, Appl
14	5224.4	99.1	5304	43	US-10-244-633-2	Sequence 2, Appl
15	4925.4	93.4	7369	75	US-60-172-360-23886	Sequence 23886, A
16	4017.4	76.2	6000	52	US-10-509-595-1	Sequence 1, Appl
17	1804.4	34.2	2800	23	US-09-366-952-1	Sequence 1, Appl
18	1804.4	34.2	2800	43	US-10-278-698-294	Sequence 294, App
19	1804.4	34.2	2800	43	US-10-278-698-808	Sequence 808, App
20	1804.4	34.2	2800	62	US-10-803-557-10	Sequence 10, Appl
21	1804.4	34.2	2800	64	US-10-956-243-1	Sequence 1, Appl
22	1691.6	32.1	2666	17	US-08-791-154-3	Sequence 3, Appl
23	1690	32.1	2397	17	US-08-791-154-1	Sequence 1, Appl

Qy	3601	CCTGATTTCTAAATCTATATATTTTCCTTTACAAGCTGAGTAATTTCTGAGCAAGTCAACG	3666
Db	3601	CCTGATTTCTAAATCTATATATTTTCCTTTACAAGCTGAGTAATTTCTGAGCAAGTCAACG	3660
Qy	3661	GTAGTAACTCAGGCTGTAAGATTACTTTAGTTTCTCCCTATTATAGGAACCTCTTTTCTCTGT	3720
Db	3661	GTAGTAACTCAGGCTGTAAGATTACTTTAGTTTCTCCCTATTATAGGAACCTCTTTTCTCTGT	3720
Qy	3721	GGAGTTAGCAGCACAAGGGCAATCCCGTTTCTTTTAAACAGGAAGAAACAATTCCTTAAGAG	3780
Db	3721	GGAGTTAGCAGCACAAGGGCAATCCCGTTTCTTTTAAACAGGAAGAAACAATTCCTTAAGAG	3780
Qy	3781	TAAAGCCAAACAGATTCAAGCCTAGGTCTTGGCTGACTATATGATGGTTTGTGTTTTGAATAAT	3840
Db	3781	TAAAGCCAAACAGATTCAAGCCTAGGTCTTGGCTGACTATATGATGGTTTGTGTTTTGAATAAT	3840
Qy	3841	CATTTCAGCGATGTTACTATCTGATTCAAGAAATGAGACTAGTACCCTTTGGTCAGCTG	3900
Db	3841	CATTTCAGCGATGTTACTATCTGATTCAAGAAATGAGACTAGTACCCTTTGGTCAGCTG	3900
Qy	3901	TAAACAACACCCAGTTGTAATGTCTCAAGTTTCAGGCTTAACTGCAGAACCAATCAAAA	3960
Db	3901	TAAACAACACCCAGTTGTAATGTCTCAAGTTTCAGGCTTAACTGCAGAACCAATCAAAA	3960
Qy	3961	AGAATGAATCTTTTAGAGCAAACTGTGTTTCTCCACATCTGGAGGTGAGTCTCCAGGGC	4020
Db	3961	AGAATGAATCTTTTAGAGCAAACTGTGTTTCTCCACATCTGGAGGTGAGTCTCCAGGGC	4020
Qy	4021	AGTTTGGAAATATTTACTTCCAAAGTATGACACTGTGTTGGTATTAACAACATAAAGT	4080
Db	4021	AGTTTGGAAATATTTACTTCCAAAGTATGACACTGTGTTGGTATTAACAACATAAAGT	4080
Qy	4081	TGCTCAAAAGCAATCATTTATTTCAAGTGGCTTAAAGTTACTCTCGACAGTTTGGTATAT	4140
Db	4081	TGCTCAAAAGCAATCATTTATTTCAAGTGGCTTAAAGTTACTCTCGACAGTTTGGTATAT	4140
Qy	4141	TTATTTGCTATTGCCATTGCTTTTGTGTTTTTCTCTTTGGGTTTATTAATGTAAGCAG	4200
Db	4141	TTATTTGCTATTGCCATTGCTTTTGTGTTTTTCTCTTTGGGTTTATTAATGTAAGCAG	4200
Qy	4201	GGATTATTAACCTACAGTCCAGAAAGCCTGTGAATTTGAATGAGGAAAAAATTTACATTTT	4260
Db	4201	GGATTATTAACCTACAGTCCAGAAAGCCTGTGAATTTGAATGAGGAAAAAATTTACATTTT	4260
Qy	4261	TGTTTTTACCACCTTCTAACTAAATTTAAACATTTTATTTCCATTGCGAATAGAGCCATAA	4320
Db	4261	TGTTTTTACCACCTTCTAACTAAATTTAAACATTTTATTTCCATTGCGAATAGAGCCATAA	4320
Qy	4321	CTCAAAGTGGTAATAACAGTACCTGTGAATTTTGTTCATTACCAATAGNAATTCAGACATT	4380
Db	4321	CTCAAAGTGGTAATAACAGTACCTGTGAATTTTGTTCATTACCAATAGNAATTCAGACATT	4380
Qy	4381	TTATACTATATTACAGTTGTTGACAGATACGTTGTGAAGTGAATAATTTATACTCAAACCTA	4440
Db	4381	TTATACTATATTACAGTTGTTGACAGATACGTTGTGAAGTGAATAATTTATACTCAAACCTA	4440
Qy	4441	CTTTGAAATTAGACCTCCTGTCGATCTTGTGTTTTTAAACATATTAATAAAACATGTTTTAA	4500
Db	4441	CTTTGAAATTAGACCTCCTGTCGATCTTGTGTTTTTAAACATATTAATAAAACATGTTTTAA	4500
Qy	4501	ATTTTGTATTTTGTGATAATCATATTTTCATTATCATTTGTTTTCCCTTGTGAATCTATATTTT	4560
Db	4501	ATTTTGTATTTTGTGATAATCATATTTTCATTATCATTTGTTTTCCCTTGTGAATCTATATTTT	4560
Qy	4561	ATATATTTGAAAAACATCTTTCTGAGAAGAGTTCCCCAGATTTTCCAAATGAGGTTCTTGG	4620
Db	4561	ATATATTTGAAAAACATCTTTCTGAGAAGAGTTCCCCAGATTTTCCAAATGAGGTTCTTGG	4620
Qy	4621	CATGCACACACAGAGTGAAGACTGATTTAGAGGCTTAACTTGCACATTTGGTCCTCGAGA	4680
Db	4621	CATGCACACACAGAGTGAAGACTGATTTAGAGGCTTAACTTGCACATTTGGTCCTCGAGA	4680
Qy	4681	TGCAAGACTGAAATTTAGAAAGTTCTCCCAAAGATACACAGTTGTTTTTAAAGCTAGGGGTG	4740

[illegible]

RESULT 2

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US-10-244-633-34
; Sequence 34, Application US/10244633
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits,
; TITLE OF INVENTION: Prognosis And Treatm
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: 07425.0057.US01
; CURRENT APPLICATION NUMBER: US/10/244, 633
; CURRENT FILING DATE: 2002-09-17
; PRIOR APPLICATION NUMBER: US/09/306, 828
; PRIOR FILING DATE: 1999-05-07
; PRIOR APPLICATION NUMBER: US 09/227, 881
; PRIOR FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 34
; LENGTH: 5271
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-244-633-34

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Query Match      100.0%; Score 5271; DB 43; Length 5271;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 5271: Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy	1	ATCTTTGTTTCAGTTTTACCTCAGGGCTATTATGAATGAAATGAGATAACCAATGTGAAG	60
<hr/>			
ph	1	ATCTTTGTTTCAGTTTTACCTCAGGGCTATTATGAATGAAATGAGATAACCAATGTGAAG	60

QY 61 TCCTATAAATGTTATAGCTCCATTCGGATGATGTCCTTGGCAGGATGATAAGAAATCA 120
DB 61 TCCTATAAATGTTATAGCTCCATTCGGATGATGTCCTTGGCAGGATGATAAGAAATCA 120
QY 121 GGAAGAAGGAGTATCCAGTTAGCCAAAGTGTCCAGGCTGTCTGCTCTTATTTAGTGA 180
DB 121 GGAAGAAGGAGTATCCAGTTAGCCAAAGTGTCCAGGCTGTCTGCTCTTATTTAGTGA 180
QY 181 CAGATGTTGCTCTCGACAGAGCTATCTTCAGGAAACATCACAATCCAAATGTAATC 240
DB 181 CAGATGTTGCTCTCGACAGAGCTATCTTCAGGAAACATCACAATCCAAATGTAATC 240
QY 241 CATCAAAACAGGAGCTTAAGAAACAGAAATGAGATGGCACTTGGCCAAAGGAAATGCCAG 300
DB 241 CATCAAAACAGGAGCTTAAGAAACAGAAATGAGATGGCACTTGGCCAAAGGAAATGCCAG 300
QY 301 GAGAGCAAAATGATGAAATAATAACTTTTCCCTTTGTTTTTAAATTCAGGAAAAAATG 360
DB 301 GAGAGCAAAATGATGAAATAATAACTTTTCCCTTTGTTTTTAAATTCAGGAAAAAATG 360
QY 361 ATGAGGACCAAAATCAATGAATAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
DB 361 ATGAGGACCAAAATCAATGAATAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
QY 421 TAATTAAGTATTTGTTCTTGGGAAGAGACCTCCATGTGAGCTTGTGGAATGGAA 480
DB 421 TAATTAAGTATTTGTTCTTGGGAAGAGACCTCCATGTGAGCTTGTGGAATGGAA 480
QY 481 AAACGTCAAAGCATGATCTGATCAGATCCCAAGTGGATTTATTTTAAATCCACAGAT 540
DB 481 AAACGTCAAAGCATGATCTGATCAGATCCCAAGTGGATTTATTTTAAATCCACAGAT 540
QY 541 GGCATCATCTGGGGAGGCAAGTTTCAGGAAGTCAATGTTAGCAAGGACATCAATTAAC 600
DB 541 GGCATCATCTGGGGAGGCAAGTTTCAGGAAGTCAATGTTAGCAAGGACATCAATTAAC 600
QY 601 AGCAAAATCAAAATTCGCAAAATGACGAGGAAATGGGACTGGGAAAGCTTTCATAAC 660
DB 601 AGCAAAATCAAAATTCGCAAAATGACGAGGAAATGGGACTGGGAAAGCTTTCATAAC 660
QY 661 AGTGATTAGGCAAGTGTGCAACACCTCCCGCTCTATACCAAGGAAACACAAA 720
DB 661 AGTGATTAGGCAAGTGTGCAACACCTCCCGCTCTATACCAAGGAAACACAAA 720
QY 721 ATTGACTGGCTAAGCTGACATTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA 780
DB 721 ATTGACTGGCTAAGCTGACATTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA 780
QY 781 GACATGTTAAAGGCAACAGAAATATGAAAACTGAGAGCAAAACAAA 840
DB 781 GACATGTTAAAGGCAACAGAAATATGAAAACTGAGAGCAAAACAAA 840
QY 841 GGGACCTGAGGCAATTTGCTTTAGGAAGGCCAGTTTCTTAAGGAATCTTAAGAAATC 900
DB 841 GGGACCTGAGGCAATTTGCTTTAGGAAGGCCAGTTTCTTAAGGAATCTTAAGAAATC 900
QY 901 TTGAAGAATCATGAATTTTAAACATTTAAGTATGATAAACAATATGCGATGCAATATCAG 960
DB 901 TTGAAGAATCATGAATTTTAAACATTTAAGTATGATAAACAATATGCGATGCAATATCAG 960
QY 961 TTTAGACATGGTCCCAATTTTAAAGTACAGGATACAGGATACAGTGTCCAGCTCC 1020
DB 961 TTTAGACATGGTCCCAATTTTAAAGTACAGGATACAGGATACAGTGTCCAGCTCC 1020
QY 1021 GGAATGTCAGAAATCATTAAGAAATCACTGTGTCCCATCTTAACATTTTTCAGAAATGATC 1080
DB 1021 GGAATGTCAGAAATCATTAAGAAATCACTGTGTCCCATCTTAACATTTTTCAGAAATGATC 1080
QY 1081 TGTATAGCCCTCACACAGGCGCGATGTGTCTGACCTACACCACTACACCCAA 1140
DB 1081 TGTATAGCCCTCACACAGGCGCGATGTGTCTGACCTACACCACTACACCCAA 1140

QY 1141 GTGCCTCAACCATTTGTTAAAGTGTATCTCAGTAGTCCCATTTACAAATGCCACTCCCC 1200
DB 1141 GTGCCTCAACCATTTGTTAAAGTGTATCTCAGTAGTCCCATTTACAAATGCCACTCCCC 1200
QY 1201 TGTGAGCCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCAATCAGATGT 1260
DB 1201 TGTGAGCCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCAATCAGATGT 1260
QY 1261 TACAGCCAGAAAGTCCCGTCCAGGAGTGTGCTGTCTTACACTACTCTGTATGCTCTAC 1320
DB 1261 TACAGCCAGAAAGTCCCGTCCAGGAGTGTGCTGTCTTACACTACTCTGTATGCTCTAC 1320
QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCCTCC 1380
DB 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCCTCC 1380
QY 1381 CGCTGAGTGGGACTACAGGCGCACCGCCGGCTAAATTTTGTATTTAGTAGAGATGGG 1440
DB 1381 CGCTGAGTGGGACTACAGGCGCACCGCCGGCTAAATTTTGTATTTAGTAGAGATGGG 1440
QY 1441 GTTTCACCATATTTAGCCCGCTGTCTGTGAATCTCTGACCTCAGGTGATCCACCACCTC 1500
DB 1441 GTTTCACCATATTTAGCCCGCTGTCTGTGAATCTCTGACCTCAGGTGATCCACCACCTC 1500
QY 1501 AGCTCTCTAAAGTGTGGGATTTACAGGATGAGTCAAGGCGCCGCGCAAGGCTCAGTGT 1560
DB 1501 AGCTCTCTAAAGTGTGGGATTTACAGGATGAGTCAAGGCGCCGCGCAAGGCTCAGTGT 1560
QY 1561 TTAATAAGGAATAAATTTGAATGGTTTAACTAAACCAACAGGAAACAGACAAAAGCTCTGA 1620
DB 1561 TTAATAAGGAATAAATTTGAATGGTTTAACTAAACCAACAGGAAACAGACAAAAGCTCTGA 1620
QY 1621 TAATTTACAGGATTTTGGGATGGGAAATGGTGGCATGAGTGCCTGCTAGTCCAGAC 1680
DB 1621 TAATTTACAGGATTTTGGGATGGGAAATGGTGGCATGAGTGCCTGCTAGTCCAGAC 1680
QY 1681 CACTGTCTCTCATCATCTTTCTCCCTCATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 1740
DB 1681 CACTGTCTCTCATCATCTTTCTCCCTCATCTCTCTCTCTCTCTCTCTCTCTCTCTCT 1740
QY 1741 CACCATGCTTTTGTGTAAAGCTTCCACATCTGTTTACTGAAATAAGAGTATACATAAATAG 1800
DB 1741 CACCATGCTTTTGTGTAAAGCTTCCACATCTGTTTACTGAAATAAGAGTATACATAAATAG 1800
QY 1801 TTCCATTTGGGGCATCTGTGTGTGTATAGGGAGGAGGCGCATACCCAGAGACTCCT 1860
DB 1801 TTCCATTTGGGGCATCTGTGTGTGTATAGGGAGGAGGCGCATACCCAGAGACTCCT 1860
QY 1861 TGAAGCCCGCGCAGAGGTTTCTCTCCAGCTGGGGGAGCCCTGCAAGCAACCCGGGTCC 1920
DB 1861 TGAAGCCCGCGCAGAGGTTTCTCTCCAGCTGGGGGAGCCCTGCAAGCAACCCGGGTCC 1920
QY 1921 TGGGTGTCTGAGCAACTGCGCAGCGGCTGCACTGTTTGTGTTTATCTCTCTCTCTCTCT 1980
DB 1921 TGGGTGTCTGAGCAACTGCGCAGCGGCTGCACTGTTTGTGTTTATCTCTCTCTCTCTCT 1980
QY 1981 GACCTGTCT 2040
DB 1981 GACCTGTCT 2040
QY 2041 TAATGAGTACTTATATCTGCCAGACACCAAGCAAAATGGTGGCAAGCAAGCTCTCTCTCT 2100
DB 2041 TAATGAGTACTTATATCTGCCAGACACCAAGCAAAATGGTGGCAAGCAAGCTCTCTCTCT 2100
QY 2101 CCTACTCTGTTGGAGTGTGACAGTCTCTCTGGAAGACGTGCAAGAAATTAATAGCCA 2160
DB 2101 CCTACTCTGTTGGAGTGTGACAGTCTCTCTGGAAGACGTGCAAGAAATTAATAGCCA 2160
QY 2161 GCCAATTAACCCAGTGTGAAAGAAATTAACCAATCTTTGAAAGAAATTTGCGC 2220
DB 2161 GCCAATTAACCCAGTGTGAAAGAAATTAACCAATCTTTGAAAGAAATTTGCGC 2220
QY 2221 AGCATCCCTTAAACAGGCGACCTCTCTAGGCGCCCTCTCTCTCTCTCTCTCTCTCTCTCT 2280

Db 2221 AGCATCCCTTAAACAGGCCACCTCCCTAGCGCCCTCTGCTCCATCTGCTGCCGAGG 2280
Qy 2281 CCCCAGAGCCGAGCTTCCAAAGCCTCTCTCCATCAGTCAACAGGCTGACGCTGCGCT 2340
Db 2281 CCCCAGAGCCGAGCTTCCAAAGCCTCTCTCCATCAGTCAACAGGCTGACGCTGCGCT 2340
Qy 2341 GCTCGCTTCCGCTGAAATCGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2400
Db 2341 GCTCGCTTCCGCTGAAATCGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2400
Qy 2401 CCAGAGAGAAATGAGAGGAAATAGTCTTAACGAGAAATCTGAGAGGAGCAAGTGTTC 2460
Db 2401 CCAGAGAGAAATGAGAGGAAATAGTCTTAACGAGAAATCTGAGAGGAGCAAGTGTTC 2460
Qy 2461 CTGAGAGGAAAGGGGCTCCAGCTCCAGGAGAAATCCAGGAGGTGGGAGCTCAGGGAG 2520
Db 2461 CTGAGAGGAAAGGGGCTCCAGCTCCAGGAGAAATCCAGGAGGTGGGAGCTCAGGGAG 2520
Qy 2521 TGGGAGCGCTGGGCTGAGCGGCTGCTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 2580
Db 2521 TGGGAGCGCTGGGCTGAGCGGCTGCTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 2580
Qy 2581 GCTGCCAGATGTCAGTGTGTTTACGGGCTGGGAGTTCCTGCTGCTGCTGCTGCTGCTG 2640
Db 2581 GCTGCCAGATGTCAGTGTGTTTACGGGCTGGGAGTTCCTGCTGCTGCTGCTGCTGCTG 2640
Qy 2641 CTTTATCTCTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2700
Db 2641 CTTTATCTCTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2700
Qy 2701 ATAAAGTCAGCTGTTAAATTCAGAGGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2760
Db 2701 ATAAAGTCAGCTGTTAAATTCAGAGGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2760
Qy 2761 TTAATGGAATATAGGAGGAGCTCAATTCCTAGGCGCTGCTGCTGCTGCTGCTGCTGCTGCT 2820
Db 2761 TTAATGGAATATAGGAGGAGCTCAATTCCTAGGCGCTGCTGCTGCTGCTGCTGCTGCTGCT 2820
Qy 2821 TGGAGTCTTTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2880
Db 2821 TGGAGTCTTTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2880
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Db 2881 TGAAGAGCTGGAAGGAGGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2940
Qy 2941 GGTGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3000
Db 2941 GGTGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3000
Qy 3001 ATAAAGACCTTGCAGCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3060
Db 3001 ATAAAGACCTTGCAGCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3060
Qy 3061 GGATGTTGAGAGGAGGAGGAGGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3120
Db 3061 GGATGTTGAGAGGAGGAGGAGGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3120
Qy 3121 GGACAGAGGAGGAGGAGGAGGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3180
Db 3121 GGACAGAGGAGGAGGAGGAGGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3180
Qy 3181 CAGGACCGAGAGGAGGAGGAGGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3240
Db 3181 CAGGACCGAGAGGAGGAGGAGGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3240
Qy 3241 TCCCTAAGCATAGACATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3300
Db 3241 TCCCTAAGCATAGACATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3300
Qy 3301 GGTAGCTTTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3360

Db 3301 GGTAGCTTTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3360
Qy 3361 TTAAACTTTTACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3420
Db 3361 TTAAACTTTTACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3420
Qy 3421 AGTGACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3480
Db 3421 AGTGACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3480
Qy 3481 ACAGATTCATCAAGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3540
Db 3481 ACAGATTCATCAAGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3540
Qy 3541 GTTCTAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3600
Db 3541 GTTCTAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 3600
Qy 3601 CTTGATTTCTAATCTAATCTAATCTAATCTAATCTAATCTAATCTAATCTAATCTAATCT 3660
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Qy 3661 GTAGTAACTGAGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3720
Db 3661 GTAGTAACTGAGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3720
Qy 3721 GGAGTTAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 3780
Db 3721 GGAGTTAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 3780
Qy 3781 TAAAGCAACAGATTCAGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3840
Db 3781 TAAAGCAACAGATTCAGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3840
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Db 3841 CATTTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3900
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Db 3901 TAAACAAACACCCAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3960
Qy 3961 AGAATAGAAATCTTTAGAGCAACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4020
Db 3961 AGAATAGAAATCTTTAGAGCAACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4020
Qy 4021 AGTTTGGAAATATTTTACCAAGTATTTGACACTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4080
Db 4021 AGTTTGGAAATATTTTACCAAGTATTTGACACTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4080
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Db 4081 TGCTCAAGGCAATCAATTTTCAAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4140
Qy 4141 TTATTTGGCTATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4200
Db 4141 TTATTTGGCTATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4200
Qy 4201 GGATTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTT 4260
Db 4201 GGATTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTT 4260
Qy 4261 TGTTTTTACCCTTCTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 4320
Db 4261 TGTTTTTACCCTTCTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 4320
Qy 4321 CTCAAGTGTGTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 4380
Db 4321 CTCAAGTGTGTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 4380
Qy 4381 TTATATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 4440
Db 4381 TTATATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 4440

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QY 4441 CTTTGAATAGACCTCTCTGCTGATCTTGTGTTTAAACATATTAATAAATGTTTAAA 4500
Db 4441 CTTTGAATAGACCTCTCTGCTGATCTTGTGTTTAAACATATTAATAAATGTTTAAA 4500
QY 4501 ATTTTGAATTTTGAATATCAATTTTCAATATCAATTTTGTGTTTCCCTTGTGTTTATATTTT 4560
Db 4501 ATTTTGAATTTTGAATATCAATTTTCAATATCAATTTTGTGTTTCCCTTGTGTTTATATTTT 4560
QY 4561 ATATATTTGAAAACATCTTCTGAGAAGAGTTCCTCCAGATTTTCAACATGAGTTCTTGG 4620
Db 4561 ATATATTTGAAAACATCTTCTGAGAAGAGTTCCTCCAGATTTTCAACATGAGTTCTTGG 4620
QY 4621 CATGCACACACAGAGTAAAGTCTGATTTAGAGGCTAAACATTCACATTTGCTGCTGAGA 4680
Db 4621 CATGCACACACAGAGTAAAGTCTGATTTAGAGGCTAAACATTCACATTTGCTGCTGAGA 4680
QY 4681 TGCAAGACTGAAATAGAAAGTTCTCCCAAGATPACACAGTGTGTTTAAAGCTAGGGGTG 4740
Db 4681 TGCAAGACTGAAATAGAAAGTTCTCCCAAGATPACACAGTGTGTTTAAAGCTAGGGGTG 4740
QY 4741 AGGGGGAAATCTCCCGCTCTATAGGAATGCTCTCCCTGGAGCTGTTAGGGTGTCTGTC 4800
Db 4741 AGGGGGAAATCTCCCGCTCTATAGGAATGCTCTCCCTGGAGCTGTTAGGGTGTCTGTC 4800
QY 4801 CTGTGTTCTGGCTGGCTGTTATTTTCTGCTCCCTGCTACGTCCTTAAAGGACTTCTTT 4860
Db 4801 CTGTGTTCTGGCTGGCTGTTATTTTCTGCTCCCTGCTACGTCCTTAAAGGACTTCTTT 4860
QY 4861 GGATCTCCAGTTCCTAGCATAGTGCCTGGACAGTGCAGGTTCTCAATGAGTTTGCAGAG 4920
Db 4861 GGATCTCCAGTTCCTAGCATAGTGCCTGGACAGTGCAGGTTCTCAATGAGTTTGCAGAG 4920
QY 4921 TGAATGGAATATTAACCTAGAAATATATCTTGTGTAATCAGACACACAGTGTCTCTGG 4980
Db 4921 TGAATGGAATATTAACCTAGAAATATATCTTGTGTAATCAGACACACAGTGTCTCTGG 4980
QY 4981 TGTAAAGTGTGTACGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 5040
Db 4981 TGTAAAGTGTGTACGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 5040
QY 5041 TAGGAATATTTATTTGGGATGATGGTGATGATGATGATGATGATGATGATGATGATGATGAT 5100
Db 5041 TAGGAATATTTATTTGGGATGATGGTGATGATGATGATGATGATGATGATGATGATGATGAT 5100
QY 5101 CAAACAGACTCTGGAAGTTATTTTCTAGAAATCTTGTGCGAGCGTGAAGGCAACCC 5160
Db 5101 CAAACAGACTCTGGAAGTTATTTTCTAGAAATCTTGTGCGAGCGTGAAGGCAACCC 5160
QY 5161 CCTGTGCACAGCCCAACCCAGCTCAGTGGCCACCTCTGCTTCTCCCAATGAAGGCTG 5220
Db 5161 CCTGTGCACAGCCCAACCCAGCTCAGTGGCCACCTCTGCTTCTCCCAATGAAGGCTG 5220
QY 5221 GCTCCCAATGATATTAACCTCTCTGAGCTCGGGCATGAGCCAGCAAGG 5271
Db 5221 GCTCCCAATGATATTAACCTCTCTGAGCTCGGGCATGAGCCAGCAAGG 5271

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RESULT 3
US-08-938-669-3
; Sequence 3, Application US/08938669
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS,
; PROGNOSIS AND TREATMENT OF GLAUCOMA AND RELATED DISEASE
; TITLE OF INVENTION: S
; NUMBER OF SEQUENCES: 32
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Howrey & Simon
; STREET: 1299 Pennsylvania Avenue, N.W.
; CITY: Washington
; STATE: DC

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; COUNTRY: USA
; ZIP: 20004-2402
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/938,669
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/791,154
; FILING DATE: 28-JAN-1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Sira, Serge
; REGISTRATION NUMBER: 39,445
; REFERENCE/DOCKET NUMBER: 07425-0034
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202 383-6857
; TELEFAX: 202 383-6610
; TELEX:
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 6169 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-08-938-669-3

Query Match 100.0%; Score 5271; DB 19; Length 6169;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 5271; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAAATGAAATGAGATAACCAATGTGAAAG 60
Db 1 ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAAATGAAATGAGATAACCAATGTGAAAG 60
QY 61 TCCTATAACTGTATAGCTCCATTCGGATGTATGCTTTTGGCAGGATGATAAGATCA 120
Db 61 TCCTATAACTGTATAGCTCCATTCGGATGTATGCTTTTGGCAGGATGATAAGATCA 120
QY 121 GGAAGAGGAGTATCCACGTTAGCCAAAGTGTCCAGGCTGTCTGCTCTTATTTAGTGA 180
Db 121 GGAAGAGGAGTATCCACGTTAGCCAAAGTGTCCAGGCTGTCTGCTCTTATTTAGTGA 180
QY 181 CAGATGTTCTCTGACAGAACTATTCTTCAGGAAACATCATCAATATGTTAAATC 240
Db 181 CAGATGTTCTCTGACAGAACTATTCTTCAGGAAACATCATCAATATGTTAAATC 240
QY 241 CATCAACAGGAGCTTAAGAAACAGGATGAGATGGGCACTTCCCAAGGAAATGCCAG 300
Db 241 CATCAACAGGAGCTTAAGAAACAGGATGAGATGGGCACTTCCCAAGGAAATGCCAG 300
QY 301 GAGAGCAAAATAATGATGAAATAATAACTTTTCCCTTTGTTTAAATTTTCAAGGAAAAATG 360
Db 301 GAGAGCAAAATAATGATGAAATAATAACTTTTCCCTTTGTTTAAATTTTCAAGGAAAAATG 360
QY 361 ATGAGGACCAAAATCAATGAATAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
Db 361 ATGAGGACCAAAATCAATGAATAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
QY 421 TAATTAAGTATTGTTTCTTGGGAGAGACCTCCATGTCAGCTTGATGGGAAATGGAA 480
Db 421 TAATTAAGTATTGTTTCTTGGGAGAGACCTCCATGTCAGCTTGATGGGAAATGGAA 480
QY 481 AAAAGTCAAAAGCATGATCTGATCAGATCCCAAGTGGATTTATTTTAAAAACCAGAT 540
Db 481 AAAAGTCAAAAGCATGATCTGATCAGATCCCAAGTGGATTTATTTTAAAAACCAGAT 540
QY 541 GGCATCCTCTCGGGAGGCAAGTTTCAGGAAAGTGTATGTTAGCAAGGACATCAACAATAC 600
Db 541 GGCATCCTCTCGGGAGGCAAGTTTCAGGAAAGTGTATGTTAGCAAGGACATCAACAATAC 600

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QY	601	AGCAAAATCAAAATTCGGCAAAATGCAGAGGAAATGGGACTGGGAAAGCTTTTCAATAC	660
DB	601	AGCAAAATCAAAATTCGGCAAAATGCAGAGGAAATGGGACTGGGAAAGCTTTTCAATAC	660
QY	661	AGTGATAGGAGTGGACCATGTTCCCAACACCTCCCGTCTATATACAGGGAACACAAA	720
DB	661	AGTGATAGGAGTGGACCATGTTCCCAACACCTCCCGTCTATATACAGGGAACACAAA	720
QY	721	ATTGACTGGGCTAAGCTGGACTTTTCAAGGGAATATGAAAACTGAGAGCAAAACAAA	780
DB	721	ATTGACTGGGCTAAGCTGGACTTTTCAAGGGAATATGAAAACTGAGAGCAAAACAAA	780
QY	781	GACATGGTTAAAGGCAACAGAAATTTGTAGCCTTTAAAGCAGAGTGCCTCAGCA	840
DB	781	GACATGGTTAAAGGCAACAGAAATTTGTAGCCTTTAAAGCAGAGTGCCTCAGCA	840
QY	841	GGGACCTGAGGCAATTTGCTTTAGGAGGCGAGTTTCTTAAGGAATCTTAAGAACTC	900
DB	841	GGGACCTGAGGCAATTTGCTTTAGGAGGCGAGTTTCTTAAGGAATCTTAAGAACTC	900
QY	901	TTGAAAGATCATGAATTTTAAACATTTTAAAGTATAAAACAAATPATCGATGCAATACAG	960
DB	901	TTGAAAGATCATGAATTTTAAACATTTTAAAGTATAAAACAAATPATCGATGCAATACAG	960
QY	961	TTTATGATGGTCCCAATTTTATATAAGTACAGGATACAGGATAAGTGTCCAGCTCC	1020
DB	961	TTTATGATGGTCCCAATTTTATATAAGTACAGGATACAGGATAAGTGTCCAGCTCC	1020
QY	1021	GGATAGTTCAGAAATCATTTAGAAATCACTGTGTCCCATCTTAATCTTTTCAAGATGATC	1080
DB	1021	GGATAGTTCAGAAATCATTTAGAAATCACTGTGTCCCATCTTAATCTTTTCAAGATGATC	1080
QY	1081	TGTATAGGCTCACACAGGCGCGATGTGTCTGACCTACACACCATCTACACCCAA	1140
DB	1081	TGTATAGGCTCACACAGGCGCGATGTGTCTGACCTACACACCATCTACACCCAA	1140
QY	1141	GTGCTCAACCATTTGTATAGTGTATCTCAGTAGTCCCATTTACAAATGCGACCTCC	1200
DB	1141	GTGCTCAACCATTTGTATAGTGTATCTCAGTAGTCCCATTTACAAATGCGACCTCC	1200
QY	1201	TGTGAGGCTCATCCGCTCCACAGGAGTCTCCCACTCTAGACTTTCTGCATCAGAGT	1260
DB	1201	TGTGAGGCTCATCCGCTCCACAGGAGTCTCCCACTCTAGACTTTCTGCATCAGAGT	1260
QY	1261	TACAGCAGAGTCTCCGCTGAGGCTGAGGCTGTGTCTTACACCTACCTGTATGCTTAC	1320
DB	1261	TACAGCAGAGTCTCCGCTGAGGCTGAGGCTGTGTCTTACACCTACCTGTATGCTTAC	1320
QY	1321	ACCTGAGCTCATCTGCAACCTCTGCTCCAGGTTCAAGGAATTTCTCTGTCTCAGCTCC	1380
DB	1321	ACCTGAGCTCATCTGCAACCTCTGCTCCAGGTTCAAGGAATTTCTCTGTCTCAGCTCC	1380
QY	1381	CGGCTAGCTGGGACTACAGGCGCACCGCGCTAAATTTTGTATTTGTATGATAGATGG	1440
DB	1381	CGGCTAGCTGGGACTACAGGCGCACCGCGCTAAATTTTGTATTTGTATGATAGATGG	1440
QY	1441	GTTTTACCATATTAGCCCGGCTGTCTTGAACCTCTGACCTCAGGTGATCCACCCCTC	1500
DB	1441	GTTTTACCATATTAGCCCGGCTGTCTTGAACCTCTGACCTCAGGTGATCCACCCCTC	1500
QY	1501	AGCCTCTTAAAGTCTGGGATTAAGGATGTTTAACTAAACCAAGGGAACAGAAAGCTGTGA	1560
DB	1501	AGCCTCTTAAAGTCTGGGATTAAGGATGTTTAACTAAACCAAGGGAACAGAAAGCTGTGA	1560
QY	1561	TTAATAGGAATTAATTTGAATGTTTAACTAAACCAAGGGAACAGAAAGCTGTGA	1620
DB	1561	TTAATAGGAATTAATTTGAATGTTTAACTAAACCAAGGGAACAGAAAGCTGTGA	1620
QY	1621	TAATTTTTCAGGATTTCTTGGATGGGAAATGGTGCATGAGCTGCCTGCTAGTCCAGAC	1680
DB	1621	TAATTTTTCAGGATTTCTTGGATGGGAAATGGTGCATGAGCTGCCTGCTAGTCCAGAC	1680

QY	1681	CACCTGCTCATCATCTTTCTTCCCTCATCTCTCATTTTCAGGCTAAGTTACCATTTTAT	1740
DB	1681	CACCTGCTCATCATCTTTCTTCCCTCATCTCTCATTTTCAGGCTAAGTTACCATTTTAT	1740
QY	1741	CACCATGCTTTTGTGTAGGCTCCACATCGTTACTGAAATAGAGTATACATAAATAG	1800
DB	1741	CACCATGCTTTTGTGTAGGCTCCACATCGTTACTGAAATAGAGTATACATAAATAG	1800
QY	1801	TTCCATTTTGGGCTCATCTGTGTGTGTATAGGAGGAGGAGGATACCCAGAGACTCT	1860
DB	1801	TTCCATTTTGGGCTCATCTGTGTGTGTATAGGAGGAGGAGGATACCCAGAGACTCT	1860
QY	1861	TGAAGCCCTCCGCGCAGAGTTTCTCTCAGCTGTGGGAGCCCTGCAAGCACCCCGGCTCC	1920
DB	1861	TGAAGCCCTCCGCGCAGAGTTTCTCTCAGCTGTGGGAGCCCTGCAAGCACCCCGGCTCC	1920
QY	1921	TGGGTGTCTGAGCAACCTGCGAGCCGCTGCGCACTGTGTGTGTGTATCATCTCTAGG	1980
DB	1921	TGGGTGTCTGAGCAACCTGCGAGCCGCTGCGCACTGTGTGTGTGTATCATCTCTAGG	1980
QY	1981	GACCTGTTGCTTTCTATTTCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT	2040
DB	1981	GACCTGTTGCTTTCTATTTCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT	2040
QY	2041	TATTGAGTACTTATATCTGCGCAGACACAGAGACAAATGTGTGTGTGTGTGTGTGTGT	2100
DB	2041	TATTGAGTACTTATATCTGCGCAGACACAGAGACAAATGTGTGTGTGTGTGTGTGTGT	2100
QY	2101	CCTACCTTGTGGAGGTGACAGTGTCTCATGGAAGACGTGTGTGTGTGTGTGTGTGTGT	2160
DB	2101	CCTACCTTGTGGAGGTGACAGTGTCTCATGGAAGACGTGTGTGTGTGTGTGTGTGTGT	2160
QY	2161	GCCAACTTAAACCCAGT	2220
DB	2161	GCCAACTTAAACCCAGT	2220
QY	2221	AGCATCTTAAACAGGCGCCTCTCTAGCGCCCTCTGTGTGTGTGTGTGTGTGTGTGTGT	2280
DB	2221	AGCATCTTAAACAGGCGCCTCTCTAGCGCCCTCTGTGTGTGTGTGTGTGTGTGTGTGT	2280
QY	2281	CCCCAAGCCCGAGTCTTCCAAAGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	2340
DB	2281	CCCCAAGCCCGAGTCTTCCAAAGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	2340
QY	2341	GCTGTCTTCCCGT	2400
DB	2341	GCTGTCTTCCCGT	2400
QY	2401	CCAGAAAGGAAATGAGAGGAGGAAATAGTCTTAAAGGAGAACTGTGTGTGTGTGTGT	2460
DB	2401	CCAGAAAGGAAATGAGAGGAGGAAATAGTCTTAAAGGAGAACTGTGTGTGTGTGTGT	2460
QY	2461	CTCAGAGGAAAGGCGCTTCCAGTCCAGAGAAATTCAGAGAGTGTGTGTGTGTGTGTGT	2520
DB	2461	CTCAGAGGAAAGGCGCTTCCAGTCCAGAGAAATTCAGAGAGTGTGTGTGTGTGTGTGT	2520
QY	2521	TGGGAGCTGTGGGCTGT	2580
DB	2521	TGGGAGCTGTGGGCTGT	2580
QY	2581	GCTGTCCAGATGT	2640
DB	2581	GCTGTCCAGATGT	2640
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DB	2641	CTTTTATCTTTTCTGT	2700
QY	2701	ATAAGTACGCTGT	2760
DB	2701	ATAAGTACGCTGT	2760
QY	2761	TTAATGGGAAATAGGAGGAGTCTATTTCTCTAGGCGCTTAAATTCACCGAAGAGTGAC	2820

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 DB 661 AGTGAATPAGCAGTTGACCAATGTTGCGCAACATCTCCCGCTCTATACAGGGAACACAAA 720
 QY 721 ATTGACTGGGCTAAGCCTGAGCTTTCAAGGGAATATGAAAACTAGAGACAAACAAA 780
 DB 721 ATTGACTGGGCTAAGCCTGAGCTTTCAAGGGAATATGAAAACTAGAGACAAACAAA 780
 QY 781 GACATGCTTAAAGGCAACAGAACATTTGTGAGCCTTCAAAGCAGCAGTCCCTCAGCA 840
 DB 781 GACATGCTTAAAGGCAACAGAACATTTGTGAGCCTTCAAAGCAGCAGTCCCTCAGCA 840
 QY 841 GGGACCTGAGGCAATTTGCTTTAGGAAGGCGAGTTTCTTAAGGAATCTTAAGAACTC 900
 DB 841 GGGACCTGAGGCAATTTGCTTTAGGAAGGCGAGTTTCTTAAGGAATCTTAAGAACTC 900
 QY 901 TTGGAAGCATGATGAATTTTAAACCAATTTTAAAGTATAAAACAAATATGCGATGCAATACAG 960
 DB 901 TTGGAAGCATGATGAATTTTAAACCAATTTTAAAGTATAAAACAAATATGCGATGCAATACAG 960
 QY 961 TTTAGACATGGGCTCCCAATTTTATAAAGTCAGGCAATCAAGGATAACGTTGCCAGTCC 1020
 DB 961 TTTAGACATGGGCTCCCAATTTTATAAAGTCAGGCAATCAAGGATAACGTTGCCAGTCC 1020
 QY 1021 GGATAGTCAGAAATCATTAAGAAATCATGTTGCCCATCTTAATCTTTTTCAGATGATC 1080
 DB 1021 GGATAGTCAGAAATCATTAAGAAATCATGTTGCCCATCTTAATCTTTTTCAGATGATC 1080
 QY 1081 TGTATAGCCCTCACACAGGCGCGATGTTCTGACCTACAAACCAATCTTACAAACCAA 1140
 DB 1081 TGTATAGCCCTCACACAGGCGCGATGTTCTGACCTACAAACCAATCTTACAAACCAA 1140
 QY 1141 GTGCTCAACCATTTGTTAAACGTTGTCATCTCAGTAGTCCCAATTAAGATGCCACTCCC 1200
 DB 1141 GTGCTCAACCATTTGTTAAACGTTGTCATCTCAGTAGTCCCAATTAAGATGCCACTCCC 1200
 QY 1201 TGTGAGCCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGATCTTGCATCAGATGT 1260
 DB 1201 TGTGAGCCCATCCCGCTCCACAGGAAGTCTCCCACTCTAGATCTTGCATCAGATGT 1260
 QY 1261 TACAGCAGAGAGCTCCGTTAGGGGTGAGGTTCTGTCTTACACCTACTCTGATGCTCTAC 1320
 DB 1261 TACAGCAGAGAGCTCCGTTAGGGGTGAGGTTCTGTCTTACACCTACTCTGATGCTCTAC 1320
 QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGTTCTCAGCTCC 1380
 DB 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGTTCTCAGCTCC 1380
 QY 1381 CGCGTAGCTGGGACTACAGGGGCGACGCCCGGCTAAATTTTGTATTTGTAGATGGG 1440
 DB 1381 CGCGTAGCTGGGACTACAGGGGCGACGCCCGGCTAAATTTTGTATTTGTAGATGGG 1440
 QY 1441 GTTTCACCATATTAGCCGGCTGGTCTTGAACCTCTGACCTCAGTGATCCACCACTC 1500
 DB 1441 GTTTCACCATATTAGCCGGCTGGTCTTGAACCTCTGACCTCAGTGATCCACCACTC 1500
 QY 1501 AGCCTCTTAAAGTGTGCGGATTTACAGGCATGATCACCGCCCGGCAAGGGTCAAGTGT 1560
 DB 1501 AGCCTCTTAAAGTGTGCGGATTTACAGGCATGATCACCGCCCGGCAAGGGTCAAGTGT 1560
 QY 1561 TTAATAAGGAATTAAGTGTGTTTAAACCAACAGGGAACAGACAAAGCTGGA 1620
 DB 1561 TTAATAAGGAATTAAGTGTGTTTAAACCAACAGGGAACAGACAAAGCTGGA 1620
 QY 1621 TAAATTCAGGGAATCTTGGGATGGGGAATGGTGCCATGAGTCTGCTAGTCCAGAC 1680
 DB 1621 TAAATTCAGGGAATCTTGGGATGGGGAATGGTGCCATGAGTCTGCTAGTCCAGAC 1680
 QY 1681 CACTGGCTCATCATCTTTCTTCCCTCATCTCTCAATTTTTCAGGCTAAGTTACATTTTAT 1740
 DB 1681 CACTGGCTCATCATCTTTCTTCCCTCATCTCTCAATTTTTCAGGCTAAGTTACATTTTAT 1740
 QY 1741 CACCATGCTTTTGTGTAAGCCTCCACATCGTTACTGAAATAGAGTATACATAAACTAG 1800

DB 1741 CACCATGCTTTTGTGTAAGCCTCCACATCGTTACTGAAATAGAGTATACATAAACTAG 1800
 QY 1801 TTCCATTTGGGCGCATCTGTGTGTGTATAGGGAGAGGGCATACCCAGAGACTCCT 1860
 DB 1801 TTCCATTTGGGCGCATCTGTGTGTGTATAGGGAGAGGGCATACCCAGAGACTCCT 1860
 QY 1861 TGAAGCCCGGCGAGAGTTTCTCTCCAGCTGGGGAGCCCTGCAAGCAACCCGGGTCC 1920
 DB 1861 TGAAGCCCGGCGAGAGTTTCTCTCCAGCTGGGGAGCCCTGCAAGCAACCCGGGTCC 1920
 QY 1921 TGGGTGTCTGAGCAACCTGCGGAGCCGTCCTGTTGTTGTTTATCACTCTCTAG 1980
 DB 1921 TGGGTGTCTGAGCAACCTGCGGAGCCGTCCTGTTGTTGTTTATCACTCTCTAG 1980
 QY 1981 GACCTGTGCTTTCTATTTCTGTGTGATCTGCTTCAATCTCAGGCAATTCATTGACAAAT 2040
 DB 1981 GACCTGTGCTTTCTATTTCTGTGTGATCTGCTTCAATCTCAGGCAATTCATTGACAAAT 2040
 QY 2041 TATTGAGTACTTATATCTCCAGACACAGAGACAAAATGCTCAGCAAAAGCAGTCACTGC 2100
 DB 2041 TATTGAGTACTTATATCTCCAGACACAGAGACAAAATGCTCAGCAAAAGCAGTCACTGC 2100
 QY 2101 CCTACCTTCTGAGAGTGACAGTTTCTCATGGAAGACGTGCAAGAAATTAATAGCCA 2160
 DB 2101 CCTACCTTCTGAGAGTGACAGTTTCTCATGGAAGACGTGCAAGAAATTAATAGCCA 2160
 QY 2161 GCCAACTTAAACCCAGTGTGAAAGAAAGAAATAAACCATCTTTGAAGAAATTTGTCGC 2220
 DB 2161 GCCAACTTAAACCCAGTGTGAAAGAAAGAAATAAACCATCTTTGAAGAAATTTGTCGC 2220
 QY 2221 AGCATCCCTTAAAGGCCACCTCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGG 2280
 DB 2221 AGCATCCCTTAAAGGCCACCTCTAGCGCCCTGCTGCTCCATCTGTCGCCGAGG 2280
 QY 2281 CCCCAGGCCGAGTCTTCAAGCCTCTCTCATCAGTCAAGCCTGTCAGCTGCGCT 2340
 DB 2281 CCCCAGGCCGAGTCTTCAAGCCTCTCTCATCAGTCAAGCCTGTCAGCTGCGCT 2340
 QY 2341 GCCTGCTTCCCGTGAATCTGCTGTCATCTGAGCTGAGACTCTTGGCTCCAGCT 2400
 DB 2341 GCCTGCTTCCCGTGAATCTGCTGTCATCTGAGCTGAGACTCTTGGCTCCAGCT 2400
 QY 2401 CCAGAAAGAAATGGAGAGGAAATAGTCTTAAAGGAGAAATCTGAGGGGACAGTGTTC 2460
 DB 2401 CCAGAAAGAAATGGAGAGGAAATAGTCTTAAAGGAGAAATCTGAGGGGACAGTGTTC 2460
 QY 2461 CTGAGGGGAAAGGGGCTCCAGTCCAGGAGAAATCCAGAGGTGGGACTGCGAGGAG 2520
 DB 2461 CTGAGGGGAAAGGGGCTCCAGTCCAGGAGAAATCCAGAGGTGGGACTGCGAGGAG 2520
 QY 2521 TGGGAGCCTGGGGCTGAGCGGGTCTGAAAGGAGGAGGTTGAAAGGGGCAAGGCTGAA 2580
 DB 2521 TGGGAGCCTGGGGCTGAGCGGGTCTGAAAGGAGGAGGTTGAAAGGGGCAAGGCTGAA 2580
 QY 2581 GCTGCCAGATGTTCAAGTGTGTTCAAGGGGCTGGGAGTTTCCGTTCTCTGTCAGC 2640
 DB 2581 GCTGCCAGATGTTCAAGTGTGTTCAAGGGGCTGGGAGTTTCCGTTCTCTGTCAGC 2640
 QY 2641 CTTTTTATCTTTCTCTGCTTGGAGGAGAAAGTCTATTTCATGAAGGGATGAGTTTC 2700
 DB 2641 CTTTTTATCTTTCTCTGCTTGGAGGAGAAAGTCTATTTCATGAAGGGATGAGTTTC 2700
 QY 2701 ATAAAGTCAGCTGTTAAATTTCCAGGGTGTGATGGGTTTCTTTCAGAGGGCTTTAT 2760
 DB 2701 ATAAAGTCAGCTGTTAAATTTCCAGGGTGTGATGGGTTTCTTTCAGAGGGCTTTAT 2760
 QY 2761 TTAATGGGAATATAGGAAGCGAGCTCAATTTCTAGGCGCTTAAATTCAGGAAGAGTGAC 2820
 DB 2761 TTAATGGGAATATAGGAAGCGAGCTCAATTTCTAGGCGCTTAAATTCAGGAAGAGTGAC 2820
 QY 2821 TGGAGTCTTTTCTTCAATGTTCTTGGGCAACTACTCAGCCCTGTGTGAGCTTGGCTTA 2880


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QY 5040 ATAGGAACATTTATTTGGGTATGGGTGCATTAATTCGGATGTTCTTTTAAAGAACT 5099
DB 5040 ATAGGAACATTTATTTGGGTATGGGTGCATTAATTCGGATGTTCTTTTAAAGAACT 5099
QY 5100 CCAACAGACTCTCGAAGCTTATTTCTAAGAACTCTGCTGCAGCGTGAAGGCAACCC 5159
DB 5100 CCAACAGACTCTCGAAGCTTATTTCTAAGAACTCTGCTGCAGCGTGAAGGCAACCC 5159
QY 5160 CCTGTGCAAGCCCCCAGCCCTCACGTGGCCACCTCTGTCTTCCCTCCATGAAGGCT 5219
DB 5160 CCTGTGCAAGCCCCCAGCCCTCACGTGGCCACCTCTGTCTTCCCTCCATGAAGGCT 5219
QY 5220 GGTCTCCAGTATATATAAACCCTCTGAGCTCGGGCATGAGCCGCAAGG 5271
DB 5220 GGTCTCCAGTATATATAAACCCTCTGAGCTCGGGCATGAGCCGCAAGG 5271

RESULT 7
US-09-227-881-1
; Sequence 1, Application US/09227881
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis, Prognosis And
; CURRENT APPLICATION NUMBER: US/09/227,881
; CURRENT FILING DATE: 1999-01-11
; EARLIER APPLICATION NUMBER: US 08/938,669
; EARLIER FILING DATE: 1997-09-26
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 1
; LENGTH: 5300
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-227-881-1

Query Match          99.5%; Score 5246.4; DB 22; Length 5300;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 5269; Conservative 0; Mismatches 1; Indels 2; Gaps 2;

QY 1 ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAATGAATGAGATAACCAATGTCAAG 60
DB 1 ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAATGAATGAGATAACCAATGTCAAG 60
QY 61 TCCTATAAATCTATAGCCTCCATTCGGATGTATGTCTTTGGCAGGATGATAAAGATCA 120
DB 61 TCCTATAAATCTATAGCCTCCATTCGGATGTATGTCTTTGGCAGGATGATAAAGATCA 120
QY 121 GGAAGAGGAGTATCCAGTTAGCCAAAGTGTCCAGGCTGTGTCTGCTTTATTTAGTGA 180
DB 121 GGAAGAGGAGTATCCAGTTAGCCAAAGTGTCCAGGCTGTGTCTGCTTTATTTAGTGA 180
QY 181 CAGATGTGCTCTGACAGAGCTATTTCTCAGAAACATCATCAATATGTTAAATC 240
DB 181 CAGATGTGCTCTGACAGAGCTATTTCTCAGAAACATCATCAATATGTTAAATC 240
QY 241 CATCAACAGGAGCTAAGAAACAGGAATGAGTGGCACTTGGCCCAAGGAAATGCCAG 300
DB 241 CATCAACAGGAGCTAAGAAACAGGAATGAGTGGCACTTGGCCCAAGGAAATGCCAG 300
QY 301 GAGAGCAAAATATGATGAAATAAATCTTTTCCCTTTGTTTAAATTTTCAGGAAAAATG 360
DB 301 GAGAGCAAAATATGATGAAATAAATCTTTTCCCTTTGTTTAAATTTTCAGGAAAAATG 360
QY 361 ATCAGGACCAAAATCAATGAATGAAGAAACAGCTCAGAAAAGATGTTTCCAAATGG 420
DB 361 ATCAGGACCAAAATCAATGAATGAAGAAACAGCTCAGAAAAGATGTTTCCAAATGG 420
QY 421 TAATTAAGTATTTGTTCTTTGGGAGAGACCTCCATGTGAGCTTGTATGGGAAATGGAA 480
DB 421 TAATTAAGTATTTGTTCTTTGGGAGAGACCTCCATGTGAGCTTGTATGGGAAATGGAA 480
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QY 481 AAAAGTCAAAAGCATGATCTGATCAGATCCCAAGTGGATTTATTTTAAAGAACT 540
DB 481 AAAAGTCAAAAGCATGATCTGATCAGATCCCAAGTGGATTTATTTTAAAGAACT 540
QY 541 GGCATCACTCTGGGAGGCAAGTTTCAGGAAGTCTATGTTAGCAAGGACATTAACAATAC 600
DB 541 GGCATCACTCTGGGAGGCAAGTTTCAGGAAGTCTATGTTAGCAAGGACATTAACAATAC 600
QY 601 AGCAAAATCAAAATTCGCAAAATGAGGAGGAAATGGGAGCTGGGAAAGCTTTTATTAAC 660
DB 601 AGCAAAATCAAAATTCGCAAAATGAGGAGGAAATGGGAGCTGGGAAAGCTTTTATTAAC 660
QY 661 AGTGATTAAGGAGTGTGACCATGTTTCGCAACACCTCCCGTCTATACCAAGGAAACAAAA 720
DB 661 AGTGATTAAGGAGTGTGACCATGTTTCGCAACACCTCCCGTCTATACCAAGGAAACAAAA 720
QY 721 ATTGACTGGGCTTAAGCTGGACCTTTCAAGGGAATATGAAAACTCGAGAGCAAAACAAA 780
DB 721 ATTGACTGGGCTTAAGCTGGACCTTTCAAGGGAATATGAAAACTCGAGAGCAAAACAAA 780
QY 781 GACATGTTTAAAGCAACCAAGCAATTTGTGAGCTTCAAGGAGGAGTGGCCCTCAGCA 840
DB 781 GACATGTTTAAAGCAACCAAGCAATTTGTGAGCTTCAAGGAGGAGTGGCCCTCAGCA 840
QY 841 GGGACCTGAGGCAATTCGCTTTAGGAGGCGCAGTTTCTTAAGGAATCTTAAGAACTC 900
DB 841 GGGACCTGAGGCAATTCGCTTTAGGAGGCGCAGTTTCTTAAGGAATCTTAAGAACTC 900
QY 901 TTGAAAGATCATGAATTTTAAACCAATTTTAAAGTATATAAAACAAATATGCGATGATCAG 960
DB 901 TTGAAAGATCATGAATTTTAAACCAATTTTAAAGTATATAAAACAAATATGCGATGATCAG 960
QY 961 TTTAGCATGGTCCCAATTTTAAAGTACAGGCAATACAGGATACGTTGCCAGTCC 1020
DB 961 TTTAGCATGGTCCCAATTTTAAAGTACAGGCAATACAGGATACGTTGCCAGTCC 1020
QY 1021 GGATAGGTCAGAAATCATTAGAAATCAGTGTGTCCTCCATCTTAACTTTTTCAGAAATGATC 1080
DB 1021 GGATAGGTCAGAAATCATTAGAAATCAGTGTGTCCTCCATCTTAACTTTTTCAGAAATGATC 1080
QY 1081 TGTCTATAGCCCTCACACACAGGCGCGATGTGTCTGACCTTACACACATCTACACCCAA 1140
DB 1081 TGTCTATAGCCCTCACACACAGGCGCGATGTGTCTGACCTTACACACATCTACACCCAA 1140
QY 1141 GTGCTCAACCATTTGTTAAACGTTCTCTCAGTAGTCCCATTAAGATGCGCCCTCC 1200
DB 1141 GTGCTCAACCATTTGTTAAACGTTCTCTCAGTAGTCCCATTAAGATGCGCCCTCC 1200
QY 1201 TGTGAGGCGCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCATCAGATGT 1260
DB 1201 TGTGAGGCGCATCCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCATCAGATGT 1260
QY 1261 TACAGCCAGAACTCCGTGAGGCTGTGTCTTACACCTACCTGATGTCTAC 1320
DB 1261 TACAGCCAGAACTCCGTGAGGCTGTGTCTTACACCTACCTGATGTCTAC 1320
QY 1321 ACTGAGTCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGTCTCAGGCTCC 1380
DB 1321 ACTGAGTCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATCTCTGTCTCAGGCTCC 1380
QY 1381 CGCGTAGCTGGGACTACAGGCGCACCGCGCTAATTTTGTATTGTTAGTAGAGATGG 1440
DB 1381 CGCGTAGCTGGGACTACAGGCGCACCGCGCTAATTTTGTATTGTTAGTAGAGATGG 1440
QY 1441 GTTTCCACCATATTAGCCCGGCTGTTTGAATCTCTGACCTCAGGTGATCCACCCACTC 1500
DB 1441 GTTTCCACCATATTAGCCCGGCTGTTTGAATCTCTGACCTCAGGTGATCCACCCACTC 1500
QY 1501 AGCTCTCTAAAGTGTGGGATTTACAGGATGATGATCCCGCGCCCGCCCAAGGTCAGTGT 1560
DB 1501 AGCTCTCTAAAGTGTGGGATTTACAGGATGATGATCCCGCGCCCGCCCAAGGTCAGTGT 1560
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Qy	1561	TTAATAAGGAATAA	CTTGAATGGTTT	TACTAAACCAACAGGGAACAGACAAAGACTGTGA	1620
Db	1561	TTAATAAGGAATAA	CTTGAATGGTTT	TACTAAACCAACAGGGAACAGACAAAGACTGTGA	1620
Qy	1621	TAAATTTACAGGAA	TTCTGGGAATGGGAATGGTGCATAGCTGCCTGCCTAGTCCACAGAC	1680	
Db	1621	TAAATTTACAGGAA	TTCTGGGAATGGGAATGGTGCATAGCTGCCTGCCTAGTCCACAGAC	1680	
Qy	1681	CACCTGGTCTCAT	CACATCTTTCTTCCCTCATCTCTCATCTTTTCAGGCTAAGTTACCATTTTATT	1740	
Db	1681	CACCTGGTCTCAT	CACATCTTTCTTCCCTCATCTCTCATCTTTTCAGGCTAAGTTACCATTTTATT	1740	
Qy	1741	CACCAATGCTTTT	TGCTGAAGCTCCACATCGTTACTGAAATAAGAGTATACATAAACTAG	1800	
Db	1741	CACCAATGCTTTT	TGCTGAAGCTCCACATCGTTACTGAAATAAGAGTATACATAAACTAG	1800	
Qy	1801	TTCCATTTTGGGGCCAT	CTGTGTGTGTATAGGGGAGAGGGGCATACCCACAGAGACTCCT	1860	
Db	1801	TTCCATTTTGGGGCCAT	CTGTGTGTGTATAGGGGAGAGGGGCATACCCACAGAGACTCCT	1860	
Qy	1861	TGAAGCCCCCGGCAGAGGTTT	CTCTCCAGCTGGGGGAGCCCTGCAAGCACCCCGGGTCC	1920	
Db	1861	TGAAGCCCCCGGCAGAGGTTT	CTCTCCAGCTGGGGGAGCCCTGCAAGCACCCCGGGTCC	1920	
Qy	1921	TGGGTGTCTGAGCAA	CTGCGACCGGTGCCACTGTGTTTGTGTTTATCACTCTCTAGG	1980	
Db	1921	TGGGTGTCTGAGCAA	CTGCGACCGGTGCCACTGTGTTTGTGTTTATCACTCTCTAGG	1980	
Qy	1981	GACCTGTTGCTTTCT	TATTTCTGTGTGACTCGTTCACTTCATCCAGGCATTCATTGACAAATT	2040	
Db	1981	GACCTGTTGCTTTCT	TATTTCTGTGTGACTCGTTCACTTCATCCAGGCATTCATTGACAAATT	2040	
Qy	2041	TATTGAGTACTTAT	TATCTGCCAGACACCCAGAGACAAAATGGTGAGCAAAAGCAGTCACTGC	2100	
Db	2041	TATTGAGTACTTAT	TATCTGCCAGACACCCAGAGACAAAATGGTGAGCAAAAGCAGTCACTGC	2100	
Qy	2101	CCTACCTTCTGTGAGGTGACA	GTGTTTCTCATGGAAAGACGTGCAGAAAGAAAATTAATAGCCA	2160	
Db	2101	CCTACCTTCTGTGAGGTGACA	GTGTTTCTCATGGAAAGACGTGCAGAAAGAAAATTAATAGCCA	2160	
Qy	2161	GCCAACTTAAACCCAGTGCT	TGAAAGAAAGGAAATAAACACCATCTTGAAGAAATTGTGCGC	2220	
Db	2161	GCCAACTTAAACCCAGTGCT	TGAAAGAAAGGAAATAAACACCATCTTGAAGAAATTGTGCGC	2220	
Qy	2221	AGCATCCCTTAAACAGGCGAC	CTCCCTAGCGCCCCCTGCTGCCTCCATCTGTGCCCGGAGG	2280	
Db	2221	AGCATCCCTTAAACAGGCGAC	CTCCCTAGCGCCCCCTGCTGCCTCCATCTGTGCCCGGAGG	2280	
Qy	2281	CCCCCAAGCCCGAGTCTTCC	AAAGCCTCTCTCCATCAGTCACAGCGCTGCAGCTGCGCT	2340	
Db	2281	CCCCCAAGCCCGAGTCTTCC	AAAGCCTCTCTCCATCAGTCACAGCGCTGCAGCTGCGCT	2340	
Qy	2341	GCCTCGCTTCCCGTGAAT	CTGCTCTGTGTCATCTGAGCTGGAGACTCTCTGGCTCCAGCT	2400	
Db	2341	GCCTCGCTTCCCGTGAAT	CTGCTCTGTGTCATCTGAGCTGGAGACTCTCTGGCTCCAGCT	2400	
Qy	2401	CCGAAAGGAAATGGAGAGG	AAACTAGTCTTAACGGAGAACTCTGAGCGGACACAGTGTTC	2460	
Db	2401	CCGAAAGGAAATGGAGAGG	AAACTAGTCTTAACGGAGAACTCTGAGCGGACACAGTGTTC	2460	
Qy	2461	CTCAGAGGAAAGGGGCT	CTCAGCTCCAGGAAATTCACAGAGGTGGGACTGCAGAGAG	2520	
Db	2461	CTCAGAGGAAAGGGGCT	CTCAGCTCCAGGAAATTCACAGAGGTGGGACTGCAGAGAG	2520	
Qy	2521	TGGGGACGCTGGGGCT	GAGCGGTGTGAAAGCAGGAAGGTGAAAAGGGCAAGGCTGAA	2580	
Db	2521	TGGGGACGCTGGGGCT	GAGCGGTGTGAAAGCAGGAAGGTGAAAAGGGCAAGGCTGAA	2580	
Qy	2581	GCTGCCAGATGTTT	CAGTGTGTTTCAAGCGGCTGGGAGTTTCCGTTGCTTCTGTGAGC	2640	
Db	2581	GCTGCCAGATGTTT	CAGTGTGTTTCAAGCGGCTGGGAGTTTCCGTTGCTTCTGTGAGC	2640	
Qy	2641	CTTTTTTATCTTTT	CTCTGCTGGAGGAAAGAACTCTATTTCATGAGGGGATGCAGTTTC	2700	

Db	2641	CTTTTATCTCTCTCTGCTGGAGGAGAAAGATCTATTTCTATGAAGGATGCAGTTTC	2700
Qy	2701	ATAAGTCAGCTGTTAAAAATCCAGGCGTGTGCATGGGTTTTCCTTCAAGAGCCCTTAT	2760
Db	2701	ATAAGTCAGCTGTTAAAAATCCAGGCGTGTGCATGGGTTTTCCTTCAAGAGCCCTTAT	2760
Qy	2761	TTAATGGGAATATAGGAAGCGAGCTCATTTTCCTAGGCCGTAAATTCACGGAGAACTGAC	2820
Db	2761	TTAATGGGAATATAGGAAGCGAGCTCATTTTCCTAGGCCGTAAATTCACGGAGAACTGAC	2820
Qy	2821	TGGAGTCCTTTTCTATGCTCTCTGGGCAACTACTCAGCCCTGTGTGGTGTGACTTGCTT	2880
Db	2821	TGGAGTCCTTTTCTATGCTCTCTGGGCAACTACTCAGCCCTGTGTGGTGTGACTTGCTT	2880
Qy	2881	TGCAGAGCGGTGCAAAACCTTGGAAACAGAGACTCGGTTTCTTCTTGGTCTGCCATT	2940
Db	2881	TGCAGAGCGGTGCAAAACCTTGGAAACAGAGACTCGGTTTCTTCTTGGTCTGCCATT	2940
Qy	2941	GGTTGGCTGTGCACCCGTGGGCAAGTGTCTCTCTTCCCTGGGCCCATAGTCTTCTCTGCT	3000
Db	2941	GGTTGGCTGTGCACCCGTGGGCAAGTGTCTCTCTTCCCTGGGCCCATAGTCTTCTCTGCT	3000
Qy	3001	ATAAGAACCTTGGCAGCTCTCTGTGTTCTGTGAAACATTTCCCTGTGTGATTTCTGTGAGGGG	3060
Db	3001	ATAAGAACCTTGGCAGCTCTCTGTGTTCTGTGAAACATTTCCCTGTGTGATTTCTGTGAGGGG	3060
Qy	3061	GGATGTTGAGGGGAGGAGGAGAGCTGGAGCAGCTGAGCCACAGGGGAGGTGAGGG	3120
Db	3061	GGATGTTGAGGGGAGGAGGAGAGCTGGAGCAGCTGAGCCACAGGGGAGGTGAGGG	3120
Qy	3121	GGACAGGAAGCGACGAGAGCTGGGTGCTCCATCAGTCTCTCACTGATCACGTCAGACTC	3180
Db	3121	GGACAGGAAGCGACGAGAGCTGGGTGCTCCATCAGTCTCTCACTGATCACGTCAGACTC	3180
Qy	3181	CAGGACCGAGAGCCAAATGCTTTCAGGAAAGCTCAATGAACCCACAGCCACATTTTCTCT	3240
Db	3181	CAGGACCGAGAGCCAAATGCTTTCAGGAAAGCTCAATGAACCCACAGCCACATTTTCTCT	3240
Qy	3241	TCCCTTAGCATGACATGCAATTTGCCAATACCAAAAGAAATGCAGAGCTAACTGGT	3300
Db	3241	TCCCTTAGCATGACATGCAATTTGCCAATACCAAAAGAAATGCAGAGCTAACTGGT	3300
Qy	3301	GGTAGCTTTTGCCTGGCATTTCAAAAATCTGGGCCAGAGCAAGTGGAAAATGCCAGAGTTG	3360
Db	3301	GGTAGCTTTTGCCTGGCATTTCAAAAATCTGGGCCAGAGCAAGTGGAAAATGCCAGAGTTG	3360
Qy	3361	TTAAACTTTTCAECCTGACACAGACCCCAACGAGCTCAGCAGTGACTGTCTGACAGCACGG	3420
Db	3361	TTAAACTTTTCAECCTGACACAGACCCCAACGAGCTCAGCAGTGACTGTCTGACAGCACGG	3420
Qy	3421	AGTGACCTGACGCGGAGGGGAGAGAAAAGAGAGGATAGTGTATGACAGAGAAAG	3480
Db	3421	AGTGACCTGACGCGGAGGGGAGAGAAAAGAGAGGATAGTGTATGACAGAGAAAG	3480
Qy	3481	ACAGATTCAATTCAAGGGCAGTGGGAATTGACCAAGGGAATTATAGTCCACGTCGCTGG	3540
Db	3481	ACAGATTCAATTCAAGGGCAGTGGGAATTGACCAAGGGAATTATAGTCCACGTCGCTGG	3540
Qy	3541	GTTCTAGGAGGCAAGGCTATATTGTGGGGGGGAAAAATCAGTTCAGGGGAAGTCCGGAGA	3600
Db	3541	GTTCTAGGAGGCAAGGCTATATTGTGGGGGGGAAAAATCAGTTCAGGGGAAGTCCGGAGA	3600
Qy	3601	CCTGATTTCTAACTACTATTTTTTCTTTTACAGCTGAGTAATTTCTGAGCAAGTCACAAG	3660
Db	3601	CCTGATTTCTAACTACTATTTTTTCTTTTACAGCTGAGTAATTTCTGAGCAAGTCACAAG	3660
Qy	3661	GTAGTAACCTGAGGCTGTAGATTTACTAGTTTCTCCTTATTAGGAACCTTTTTTCTCTGT	3720
Db	3661	GTAGTAACCTGAGGCTGTAGATTTACTAGTTTCTCCTTATTAGGAACCTTTTTTCTCTGT	3720
Qy	3721	GGAGTTAGCAGCAAAAGGGCAATCCCGTTTCTTTTAAACAGGAGAGAAAAATTTCTTCAAGAG	3780

3721	GGAGTTAGCAGCA	CAAGGGCAATCCC	GTTCCTTTTAA	CAGGAAGAAAA	CATTCCTTAAGAG	3780
3781	TAAGGCAAA	CACAGATTCAAG	CCCTAGGTCCTG	CTGCACTATATGAT	TGGTTTTTTTGA	3840
3781	TAAGGCAAA	CACAGATTCAAG	CCCTAGGTCCTG	CTGCACTATATGAT	TGGTTTTTTTGA	3840
3841	CAITTCAGCGA	TGTTACTATCTG	ATTCTGATTCAGA	AAATGAGACTAGT	ACCTTTGGTTCAG	3900
3841	CAITTCAGCGA	TGTTACTATCTG	ATTCTGATTCAGA	AAATGAGACTAGT	ACCTTTGGTTCAG	3900
3901	TAAACAAA	CACCAGTGTTG	TAATGTCTCAAG	TTCAGGCTTAA	CTGCAGAACCAAT	3959
3901	TAAACAAA	CACCAGTGTTG	TAATGTCTCAAG	TTCAGGCTTAA	CTGCAGAACCAAT	3960
3960	AAGAA	TAGAATCTTTAG	AGCAAACTGTGTT	TCTCCACATCTG	GAGGTGAGTCTGC	4019
3961	AAGAA	TAGAATCTTTAG	AGCAAACTGTGTT	TCTCCAC-TC	TGAGGTTGAGTCTGC	4019
4020	CAGTTTGGAA	ATATTTACTT	CACAAGTATTG	ACA	CTGTGTTGATTA	4079
4020	CAGTTTGGAA	ATATTTACTT	CACAAGTATTG	ACA	CTGTGTTGATTA	4079
4080	TTGCTCAAA	GGCAATCATAT	TTTCAAGTGCCT	TAAAGTTACTT	CTGCAGAGTTT	4139
4080	TTGCTCAAA	GGCAATCATAT	TTTCAAGTGCCT	TAAAGTTACTT	CTGCAGAGTTT	4139
4140	TTTATTGGCT	ATTGCCATTG	CTTTTGTGTTTT	TCTCTTTGGG	TTTTATTAA	4199
4140	TTTATTGGCT	ATTGCCATTG	CTTTTGTGTTTT	TCTCTTTGGG	TTTTATTAA	4199
4200	GGGATTATTA	ACCTTACAGT	CCAGAAAGCC	TGTGAA	TTTGAATGAGGAAAA	4259
4200	GGGATTATTA	ACCTTACAGT	CCAGAAAGCC	TGTGAA	TTTGAATGAGGAAAA	4259
4260	TTGTTTTTA	CAACCTCTTAA	CTAAATTTAA	CACTTTATTC	CAATGCGAATAG	4319
4260	TTGTTTTTA	CAACCTCTTAA	CTAAATTTAA	CACTTTATTC	CAATGCGAATAG	4319
4320	ACTCAAGT	CGGTAATACAG	TACTGATTTG	TGTCATTA	CAATAGAAATCAC	4379
4320	ACTCAAGT	CGGTAATACAG	TACTGATTTG	TGTCATTA	CAATAGAAATCAC	4379
4380	TTTATATCT	ATATTTACAG	TGTGTCAGAT	ACGTTGTA	AGTGAATTTAT	4439
4380	TTTATATCT	ATATTTACAG	TGTGTCAGAT	ACGTTGTA	AGTGAATTTAT	4439
4440	ACTTTGAAA	TTAGACCTCT	CTGCTGGATCT	TTTTTAA	CAATTAATAAAC	4499
4440	ACTTTGAAA	TTAGACCTCT	CTGCTGGATCT	TTTTTAA	CAATTAATAAAC	4499
4500	AAITTTGAT	ATTTTTGATA	TAATTCATAT	CAATTTGTTT	CTCTTGTGTA	4559
4500	AAITTTGAT	ATTTTTGATA	TAATTCATAT	CAATTTGTTT	CTCTTGTGTA	4559
4560	TATATTTT	TGAAAACTCTT	CTCAGAAAG	AGTTC	CCAGATTTTCA	4619
4560	TATATTTT	TGAAAACTCTT	CTCAGAAAG	AGTTC	CCAGATTTTCA	4619
4620	GCATGCA	CACACACAG	AGTAAAGT	CTGATTTAG	AGGTTAA	4679
4620	GCATGCA	CACACACAG	AGTAAAGT	CTGATTTAG	AGGTTAA	4679
4680	ATGCAAG	ACTGAAAT	TAGAAAGTTCT	CCCAAGAT	ACACAGTTGTT	4739
4680	ATGCAAG	ACTGAAAT	TAGAAAGTTCT	CCCAAGAT	ACACAGTTGTT	4739
4740	GAGGGGG	GAATCTG	CCCTTATAG	GAATGCTCT	CCCTGGAG	4799
4740	GAGGGGG	GAATCTG	CCCTTATAG	GAATGCTCT	CCCTGGAG	4799
4800	CTTTGTGTT	CTGGCTGG	CTGTATTTTT	CTCTGT	CCCTGTAC	4859
4800	CTTTGTGTT	CTGGCTGG	CTGTATTTTT	CTCTGT	CCCTGTAC	4859

RESULT 8

RESULTS
US-09-985-637A-1

US-09-9837A-I
: Sequence 1. Application US/09985637A

; SEQUENCE 1, APPLICATION:
: GENERAL INFORMATION:

APPLICANT: polansky. Jon

APPLICANT: POJANSKY, JOH

;; TITLE OF INVENTION: METHODS TO SCREEN AN
: TITLE OF INVENTION: TO DEVELOP GLAUCOMA

10 D
NOTION: 10 D
TITLE OF INVENTION: 10 D
13587 296
REFERENCE: 13587 296

FILE REFERENCE: I3587.296
CURRENT APPLICATION NUMBER: IIS/09/985.6377A

; CURRENT APPLICATION NUMBER: US/0
 : CURRENT FILING DATE: 2001-11-05

; CURRENT FILING DATE: 20
 - NUMBER OF SEQ TO NOS: 21

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; NUMBER OF SEQ ID NOS: 21
SOFTWARE: Blast in version 3.1

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; SOFTWARE: Pat
; CEO ID NO 1

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; SEQ ID NO 1

; LENGTH: 5300

TYPE: DNA

; ORGANISM: HO

Query Match

Query Match	99.98%	Pred. No. 0:
Best Local Similarity	99.98%	Pred. No. 0:

BEST LOCAL SIMILARITY 99.5%; FREQ: NO: 0;
Matches 5269. Conservative 0: Mismatches 1: Indels 2: Gaps 2;

1 ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAAATGAAATGAGATAACCAATGTGAAAG 60

Db 1 ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAAATGAAATGAGATAACCAATGTGAAAG 60

QY
61 TCC TATAA C T G T A T A G C C T C C A T T C G G A T G T A T G T C T T T G G C A G G A T G A T A A G A A T C A 12

Db 61 TCCTATAA ACTGTATAGCCCTCCATTCGGATGTAATGCTTTGGCAGGAIGATAAAGAAATCA 12

QY IZI GGAAAGAGGAGTATCCACGTTAGCCATAGTGTCCAGGCTGTGTCTATTATTTTGTGCTG

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[illegible]

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301 CACGCCAATATATCATATGCAAAAATAAACCTTTTCCCCCTTTGTTTATAATTTCAGGAAAAAATG 36

	Db	301	GAGAGCAAATAATGATGAATAAATAAACTTTTCCCCTTGTTTTTAATTTCACGGAAAAATG	360
	Qy	361	ATGAGGACCAAAATCAATGAATAAGAAAAACAGCTCAGAAAAAGAATGTTCCAAATTTGG	420
	Db	361	ATGAGGACCAAAATCAATGAATAAGAAAAACAGCTCAGAAAAAGAATGTTCCAAATTTGG	420
	Qy	421	TAAATTAAGTATTTGTTCCTTTGGGAAGAGACTCCCATGTGAGCTTGATCGGAAAAATGGGAA	480
	Db	421	TAAATTAAGTATTTGTTCCTTTGGGAAGAGACTCCCATGTGAGCTTGATCGGAAAAATGGGAA	480
	Qy	481	AAACGTCAAAACGCATGCTGATCAGATCCCAAGTGGATTAATTTATTTTAAAAACCAGAT	540
	Db	481	AAACGTCAAAACGCATGATCAGATCCCAAGTGGATTAATTTATTTTAAAAACCAGAT	540
	Qy	541	GGCATCACTCTCGGGAGGCAAGTTTCAGGAAGTGCATGTTAGCAAAAGGCATAAACAATAAC	600
	Db	541	GGCATCACTCTCGGGAGGCAAGTTTCAGGAAGTGCATGTTAGCAAAAGGCATTAACAATAAC	600
	Qy	601	AGCAAAATCAAAATTCGCGAAATGACGAGGAAAAATGGGACTGGGAAAAGCTTTTCATAAC	660
	Db	601	AGCAAAATCAAAATTCGCGAAATGACGAGGAAAAATGGGACTGGGAAAAGCTTTTCATAAC	660
	Qy	661	AGTGATTAAGGCAGTTTGACCATGTTTCGCAACACCTCCCCGTCTATACAGAGGAACACAAAA	720
	Db	661	AGTGATTAAGGCAGTTTGACCATGTTTCGCAACACCTCCCCGTCTATACAGAGGAACACAAAA	720
	Qy	721	ATTGACTGGGCTTAAGCTGGACTTTTCAAGGGGAAATATCAAAAACTGAGAGCAAAACAAA	780
	Db	721	ATTGACTGGGCTTAAGCTGGACTTTTCAAGGGGAAATATGAAAACTGAGAGCAAAACAAA	780
	Qy	781	GACATGGTTAAAGGCCAACAGAAATTTGTGAGCCCTTCAAAAGCAGCAGTGCCCTTCAGCA	840
	Db	781	GACATGGTTAAAGGCCAACAGAAATTTGTGAGCCCTTCAAAAGCAGCAGTGCCCTTCAGCA	840
	Qy	841	GGGACCTTGAGGCATTTGCTTTTAGGAAGGCCAGTTTTCTTTAAGGAATCTTTAAGAAATCT	900
	Db	841	GGGACCTTGAGGCATTTGCTTTTAGGAAGGCCAGTTTTCTTTAAGGAATCTTTAAGAAATCT	900
	Qy	901	TTGAAAGATCATGAATTTTAAACCATTTTAAGTATATAAAACAATATGCGATGCATAATCAC	960
	Db	901	TTGAAAGATCATGAATTTTAAACCATTTTAAGTATATAAAACAATATGCGATGCATAATCAC	960
	Qy	961	TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAAGGATAAAGTGTCCTCCAGTGCC	1020
	Db	961	TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAAGGATAAAGTGTCCTCCAGTGCC	1020
	Qy	1021	GGATAGGTCAGAAATCATAGAAATCACTGTGTCCCATCCCTAACTTTTTCAGAAATATCT	1080
	Db	1021	GGATAGGTCAGAAATCATAGAAATCACTGTGTCCCATCCCTAACTTTTTCAGAAATATCT	1080
	Qy	1081	TGTCATAGCCCTCACACACAGGCCGATGTCGTGACTCAACCAACATCTCAACGCCAA	1140
	Db	1081	TGTCATAGCCCTCACACACAGGCCGATGTCGTGACTCAACCAACATCTCAACGCCAA	1140
	Qy	1141	GTGCTCTCAACCATTTGTAAACGTGTCACTCAGTAGGTGCCCATTAACAATGCGACCTGCC	1200
	Db	1141	GTGCTCTCAACCATTTGTAAACGTGTCACTCAGTAGGTGCCCATTAACAATGCGACCTGCC	1200
	Qy	1201	TGTGAGGCCCATTCGCGTCCACAGGAAGTGTCCCCACTCTAGACTTCTGCAATCAAGATGT	1260
	Db	1201	TGTGAGGCCCATTCGCGTCCACAGGAAGTGTCCCCACTCTAGACTTCTGCAATCAAGATGT	1260
	Qy	1261	TACAGCCAGAGCTCCGTGAGGGGTGAGGGTCTGTGTCTTACACTACTGTATGCTCTAC	1320
	Db	1261	TACAGCCAGAGCTCCGTGAGGGGTGAGGGTCTGTGTCTTACACTACTGTATGCTCTAC	1320
	Qy	1321	ACCTGAGCTCACTGCAACCTCTGCGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGGCTCC	1380
	Db	1321	ACCTGAGCTCACTGCAACCTCTGCGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGGCTCC	1380
	Qy	1381	CGCGTAGCTGGGACTACAGCGCGCACGCCCGGCTAATTTTTGTATTTGTAGTAGAGTGGG	1440

Db	1381	CGCGTAGCTGGGACTACAGCGCGACGCCGCGCTAAATTTTGTATTGTTAGTAGAGATGGG	1441
Qy	1441	GTTTACCAATATTAGCCCGCGCTGGTCTTTGAACCTCTGACCTCAGGTGATCCACCCACCTC	1500
Db	1441	GTTCACCATATTAGCCCGCGCTGGTCTTTGAACCTCTGACCTCAGGTGATCCACCCACCTC	1500
Qy	1501	AGCCTCCTAAAGTGCTGGGATTACAGGCATGAGTCACCGCGCCCGCCAGGGTCAGTGT	1560
Db	1501	AGCCTCCTAAAGTGCTGGGATTACAGGCATGAGTCACCGCGCCCGCCAGGGTCAGTGT	1560
Qy	1561	TTAATAAGGAATAACTTTGAATGGTTTACTATAA CCAACAGGGAACAAGCAAAAAGCTGTGA	1620
Db	1561	TTAATAAGGAATAACTTTGAATGGTTTACTATAA CCAACAGGGAACAAGCAAAAAGCTGTGA	1620
Qy	1621	TAATTTCAGGGATTCTTTGGATGGGAAATGGTGCATGAGCTGCTGCTAGTCCAGAC	1680
Db	1621	TAATTTCAGGGATTCTTTGGATGGGAAATGGTGCATGAGCTGCTGCTAGTCCAGAC	1680
Qy	1681	CACTGCTCTCATCACTTTCTCCCTCATCCTCATTTTTCAGGCTAAGTTACCATTTTATT	1740
Db	1681	CACTGCTCTCATCACTTTCTCCCTCATCCTCATTTTTCAGGCTAAGTTACCATTTTATT	1740
Qy	1741	CACCATGCTTTTGTGTAGACCTCCACATCTGTTACTGAAATAAGAGTATACATAAACTAG	1800
Db	1741	CACCATGCTTTTGTGTAGACCTCCACATCTGTTACTGAAATAAGAGTATACATAAACTAG	1800
Qy	1801	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGGAGGGGCATACCCAGAGACTCCT	1860
Db	1801	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGGAGGGGCATACCCAGAGACTCCT	1860
Qy	1861	TGAAGCCCCCGGCAGAGGTTTCCTCTCAGCTGGGGGAGCCCTGCAAGACACCCGGGGTCC	1920
Db	1861	TGAAGCCCCCGGCAGAGGTTTCCTCTCAGCTGGGGGAGCCCTGCAAGACACCCGGGGTCC	1920
Qy	1921	TGGGTGTCTGAGCAACCTGCCAGCCGTGCGCATCTGTTGTTTGTATTACTCTTAGG	1980
Db	1921	TGGGTGTCTGAGCAACCTGCCAGCCGTGCGCATCTGTTGTTTGTATTACTCTTAGG	1980
Qy	1981	GACCTGTGCTTTCTATTCTGTGTGACTCGTTTCAATTCACAGGCATTCATTGCACAAT	2040
Db	1981	GACCTGTGCTTTCTATTCTGTGTGACTCGTTTCAATTCACAGGCATTCATTGCACAAT	2040
Qy	2041	TATTGAGTACTTATATCTGCCAGACACAGAGACAAAATGGTGAGCAAAAGCAGTCACCTGC	2100
Db	2041	TATTGAGTACTTATATCTGCCAGACACAGAGACAAAATGGTGAGCAAAAGCAGTCACCTGC	2100
Qy	2101	CCTACCTTCGTGGAGGTGACAGTTTCTATGGAAGACGTGCAGAGAAAATTAATAGCCA	2160
Db	2101	CCTACCTTCGTGGAGGTGACAGTTTCTATGGAAGACGTGCAGAGAAAATTAATAGCCA	2160
Qy	2161	GCCAACTTAAACCCAGTGCTGAAAGAAAGGAATAACCACTTTGAAGAAATGTGGCG	2220
Db	2161	GCCAACTTAAACCCAGTGCTGAAAGAAAGGAATAACCACTTTGAAGAAATGTGTGGCG	2220
Qy	2221	AGCATCCCTTAAACAGGCCACCTCCTTAGCGCCCTGCTGCTCATCTGTCGCCGAGG	2280
Db	2221	AGCATCCCTTAAACAGGCCACCTCCTTAGCGCCCTGCTGCTCATCTGTCGCCGAGG	2280
Qy	2281	CCCCAAGCCGAGTCTTCCAAAGCCTCTCTCCATCAGTCACAGCGCTCAGGTGGCCT	2340
Db	2281	CCCCAAGCCGAGTCTTCCAAAGCCTCTCTCCATCAGTCACAGCGCTCAGGTGGCCT	2340
Qy	2341	GCCTCGCTTCCGTTGAATCGTCTGCTGTCATCTGAGCTGGAGACTCCTTGGCTCCAGGCT	2400
Db	2341	GCCTCGCTTCCGTTGAATCGTCTGCTGTCATCTGAGCTGGAGACTCCTTGGCTCCAGGCT	2400
Qy	2401	CCAGAAAGGAAATGGAGAGGGAAAATAGTCTAAACGAGAAATCTGGAGGGGACAGTGTTC	2460
Db	2401	CCAGAAAGGAAATGGAGAGGGAAAATAGTCTAAACGAGAAATCTGGAGGGGACAGTGTTC	2460
Qy	2461	CTCAGAGGAAAGGGGCTTCCAGTCTAAACGAGAAATTCAGGAGGTCGGGACCTCAGGGAG	2520
Db	2461	CTCAGAGGAAAGGGGCTTCCAGTCTAAACGAGAAATTCAGGAGGTCGGGACCTCAGGGAG	2520

QY	2521	TGGGACGCTGGGCTGAGCGGCTGCTGAAAGCGCAGGAAGGTGAAAGGGCAGGCGTGA	2580
DB	2521	TGGGACGCTGGGCTGAGCGGCTGCTGAAAGCGCAGGAAGGTGAAAGGGCAGGCGTGA	2580
QY	2581	GCTGCCAGATGTTCAAGTGTGTTTCAAGCGGCTGGAGTTTTCGTTGCTTCTCTGAGC	2640
DB	2581	GCTGCCAGATGTTCAAGTGTGTTTCAAGCGGCTGGAGTTTTCGTTGCTTCTCTGAGC	2640
QY	2641	CTTTTATCTTTCTCTGCTGGAGGAGGAAGTCTATTTCATGAAGGGATGCAAGTTTC	2700
DB	2641	CTTTTATCTTTCTCTGCTGGAGGAGGAAGTCTATTTCATGAAGGGATGCAAGTTTC	2700
QY	2701	ATAAAGTCAGCTGTTTAAATTCAGGGTGTGATGGGTTTTCCTTCAAGAGGCTTTAT	2760
DB	2701	ATAAAGTCAGCTGTTTAAATTCAGGGTGTGATGGGTTTTCCTTCAAGAGGCTTTAT	2760
QY	2761	TTAATGGGAATATAGGAAGGAGCTCATTTCTAGGCCGTAAATTCACGGAAGAGTGAC	2820
DB	2761	TTAATGGGAATATAGGAAGGAGCTCATTTCTAGGCCGTAAATTCACGGAAGAGTGAC	2820
QY	2821	TGGAGTCTTTTCTTTTCATGCTCTTCTGGGCAACTACTCAGCCCTGTGGTGGACTTGGCTTA	2880
DB	2821	TGGAGTCTTTTCTTTTCATGCTCTTCTGGGCAACTACTCAGCCCTGTGGTGGACTTGGCTTA	2880
QY	2881	TGCAAGACGCTCGAAACCTTGGAAATCAGGAGCTCGGTTTCTTCTGGTCTGCGCAT	2940
DB	2881	TGCAAGACGCTCGAAACCTTGGAAATCAGGAGCTCGGTTTCTTCTGGTCTGCGCAT	2940
QY	2941	GCTTGGCTGTGGACCGTGGCAAGTCTCTCTTCTCCCTGGGCCATAGTCTTCTGCT	3000
DB	2941	GCTTGGCTGTGGACCGTGGCAAGTCTCTCTTCTCCCTGGGCCATAGTCTTCTGCT	3000
QY	3001	ATAAGACCTTTCAGCTCTCGTGTCTCTGTAACACTTCCCTGTGATTTCTCTGAGGG	3060
DB	3001	ATAAGACCTTTCAGCTCTCGTGTCTCTGTAACACTTCCCTGTGATTTCTCTGAGGG	3060
QY	3061	GGATGTTGAGAGGGAAGGAGCAGCTGAGCAGCTGAGCCACAGGGGAGGTGGAAGG	3120
DB	3061	GGATGTTGAGAGGGAAGGAGCAGCTGAGCAGCTGAGCCACAGGGGAGGTGGAAGG	3120
QY	3121	GGACAGGAGGAGGAGGAGCTGGGTCTCCATCAGTCTCTCAGTCACTGATCAGCTCAGATC	3180
DB	3121	GGACAGGAGGAGGAGGAGCTGGGTCTCCATCAGTCTCTCAGTCACTGATCAGCTCAGATC	3180
QY	3181	CAGGACCGAGAGCCAAATGCTTTCAGGAAAGCTCAATGAACCCAAACAGCCACATTTTCT	3240
DB	3181	CAGGACCGAGAGCCAAATGCTTTCAGGAAAGCTCAATGAACCCAAACAGCCACATTTTCT	3240
QY	3241	TCCCTAAGCATAGACAAATGGCATTTGCGCAATAACCAAAAGAAATGAGAGACTAAGTGT	3300
DB	3241	TCCCTAAGCATAGACAAATGGCATTTGCGCAATAACCAAAAGAAATGAGAGACTAAGTGT	3300
QY	3301	GGTAGCTTTTGGCTGGCATTTCAAAACTGGGCCAGAGCAAGTGGAAATGCCAGAGTTG	3360
DB	3301	GGTAGCTTTTGGCTGGCATTTCAAAACTGGGCCAGAGCAAGTGGAAATGCCAGAGTTG	3360
QY	3361	TTAAACTTTTTCACCTGACAGCAGCCACCCAGCAGCTCAGCAGTGAAGTGAAGCAGCAG	3420
DB	3361	TTAAACTTTTTCACCTGACAGCAGCCACCCAGCAGCTCAGCAGTGAAGTGAAGCAGCAG	3420
QY	3421	AGTGAAGCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	3480
DB	3421	AGTGAAGCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	3480
QY	3481	ACAGATTCATTCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	3540
DB	3481	ACAGATTCATTCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	3540
QY	3541	GTTCTAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	3600
DB	3541	GTTCTAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	3600

QY	3601	CCTGATTTCTAATACTATATATTTTCTTTTACAGCTGAGTAATTTCTGAGCAAGTCACAAG	3660
DB	3601	CCTGATTTCTAATACTATATATTTTCTTTTACAGCTGAGTAATTTCTGAGCAAGTCACAAG	3660
QY	3661	GTAGTAACGTAGGCTGTAAGATTAATTTAGTTTCTCTTTATTAGGAACCTCTTTTCTCTGT	3720
DB	3661	GTAGTAACGTAGGCTGTAAGATTAATTTAGTTTCTCTTTATTAGGAACCTCTTTTCTCTGT	3720
QY	3721	GGAGTTAGCAGCAGCAGGCGAATCCCGTTCTTTTAAAGGAGGAGGAGGAGGAGGAGG	3780
DB	3721	GGAGTTAGCAGCAGCAGGCGAATCCCGTTCTTTTAAAGGAGGAGGAGGAGGAGGAGG	3780
QY	3781	TAAAGCCAAACAGAGATTCAAGCTTAGTCTGCTGACTATATGATTTGTTTTTGAAGAAAT	3840
DB	3781	TAAAGCCAAACAGAGATTCAAGCTTAGTCTGCTGACTATATGATTTGTTTTTGAAGAAAT	3840
QY	3841	CATTTTCAGCGATGTTTACTATCTGATTCAGAAATGAGACTAGTAGTACCCCTTTGGTCAGCTG	3900
DB	3841	CATTTTCAGCGATGTTTACTATCTGATTCAGAAATGAGACTAGTAGTACCCCTTTGGTCAGCTG	3900
QY	3901	TAAACAAACACCCAGTTGTAATGTCTCAAGTTCAAGGCTTAACCTGCAGAACCAATCAAA	3959
DB	3901	TAAACAAACACCCAGTTGTAATGTCTCAAGTTCAAGGCTTAACCTGCAGAACCAATCAAA	3960
QY	3960	AAGATAGAATCTTTAGAGCAAACTGTGTTCTCCACATCTGAGGCTGAGTCTGCCAGGG	4019
DB	3961	AAGATAGAATCTTTAGAGCAAACTGTGTTCTCCAC-TCTGAGGCTGAGTCTGCCAGGG	4019
QY	4020	CAGTTTGGAAATATTACTTCAAGTAATGACACTGTGTTGTTGTTATTAAACAACATAAG	4079
DB	4020	CAGTTTGGAAATATTACTTCAAGTAATGACACTGTGTTGTTGTTATTAAACAACATAAG	4079
QY	4080	TTGCTCAAAGGCAATCATTTTCAAGTGGCTTAAAGTTACTTCTGACAGTTTTCGTATA	4139
DB	4080	TTGCTCAAAGGCAATCATTTTCAAGTGGCTTAAAGTTACTTCTGACAGTTTTCGTATA	4139
QY	4140	TTTATTGGCTATTGCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTT	4199
DB	4140	TTTATTGGCTATTGCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTT	4199
QY	4200	GGGATTTATAACCTACAGTCCAGAAAGCTGTGAATTTGAAATGAGGAGGAGGAGGAGG	4259
DB	4200	GGGATTTATAACCTACAGTCCAGAAAGCTGTGAATTTGAAATGAGGAGGAGGAGGAGG	4259
QY	4260	TTGTTTTTACCACCTTCTAACTAAATTTAACTTTTATTTCCATTCGATAGAGCCATAA	4319
DB	4260	TTGTTTTTACCACCTTCTAACTAAATTTAACTTTTATTTCCATTCGATAGAGCCATAA	4319
QY	4320	ACTCAAAGTGGTAAATACAGTACCTGTGATTTTGTTCATTTCCCAATAGAGGAGGAGG	4379
DB	4320	ACTCAAAGTGGTAAATACAGTACCTGTGATTTTGTTCATTTCCCAATAGAGGAGGAGG	4379
QY	4380	TTTATATCTATATTAACAGTTTGTGAGTACGTTGTAAGTGAATTTTATCTCAAAACT	4439
DB	4380	TTTATATCTATATTAACAGTTTGTGAGTACGTTGTAAGTGAATTTTATCTCAAAACT	4439
QY	4440	ACTTTGAAATTAGACCTCTGCTGGATCTTGTGTTTTTAACTAATTAATAAAGATTTAA	4499
DB	4440	ACTTTGAAATTAGACCTCTGCTGGATCTTGTGTTTTTAACTAATTAATAAAGATTTAA	4499
QY	4500	AAATTTGATTTTTCATTAATCATATTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTT	4559
DB	4500	AAATTTGATTTTTCATTAATCATATTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTT	4559
QY	4560	TATATATTGAAAAATCTTTCTGAGAGAGTTCCCGAGATTTTCAACATAGAGTTCTTGG	4619
DB	4560	TATATATTGAAAAATCTTTCTGAGAGAGTTCCCGAGATTTTCAACATAGAGTTCTTGG	4619
QY	4620	GCATGCACACACAGAGTAAGAACTGATTTAGAGGCTAACATTCGATTTGCTGCTGAG	4679
DB	4620	GCATGCACACACAGAGTAAGAACTGATTTAGAGGCTAACATTCGATTTGCTGCTGAG	4679
QY	4680	ATGCAAGACTGAAATTTAGAAAGTTCTCCCAAGATACACAGTTGTTTAAAGCTAGGGT	4739


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Db 4680 ATGCAAGACTGAAATAGAAAGTCTCCAAAGATACACAGTTGTTTAAAGCTAGGGGT 4739
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Db 4740 GAGGGGGAATATCGCCCTCTATAGGAATGCTCTCCCTGGAGCCTGGTAGGTGCTGT 4799
Qy 4800 CTTGTGTTCTGGGCTGGCTGTTATTTTCTCTGTCCCTGCTACGTCCTTAAAGGACTTGT 4859
Db 4800 CTTGTGTTCTGGGCTGGCTGTTATTTTCTCTGTCCCTGCTACGTCCTTAAAGGACTTGT 4859
Qy 4860 TGGATCTCCAGTCTCTAGCATAGTCCTGGGCAAGTGCAGTCTCAATGATTTGCGAGA 4919
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Qy 4920 GTGAATGGAATATAAAGTAAATATATATCTCTGTTGAAATCAGCACACCAAGTGTCTG 4979
Db 4920 GTGAATGGAATATAAAGTAAATATATATCTCTGTTGAAATCAGCACACCAAGTGTCTG 4979
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Db 4980 GTGTAAGTGTGTACGTTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 5039
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Db 5040 ATAGGAATATATTTGGGATATGGGTGATATAATTTGGGATGTTCTTTTAAAGAACT 5099
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Db 5100 CCAACAGACTCTGGAGGTTATTTCTAGAAATCTGCTGGGAGCTGTGAGGCAACCC 5159
Qy 5160 CCCTGTGCACAGCCCAACCCAGCTCAGCTGGGCAACCTCTGTCTTCCCATGAAGGCT 5219
Db 5160 CCCTGTGCACAGCCCAACCCAGCTCAGCTGGGCAACCTCTGTCTTCCCATGAAGGCT 5219
Qy 5220 GGCTCCCAATATATAAACCCTCTGGAGCTCGGCAATGAGCCAGCAAGG 5271
Db 5220 GGCTCCCAATATATAAACCCTCTGGAGCTCGGCAATGAGCCAGCAAGG 5271

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RESULT 9

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US-10-244-633-1
; Sequence 1, Application US/10244633
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis,
; TITLE OF INVENTION: Prognosis And Treatment Of Glaucoma And Related
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: 07425.0057.US01
; CURRENT APPLICATION NUMBER: US/10/244,633
; CURRENT FILING DATE: 2002-09-17
; PRIOR APPLICATION NUMBER: US/09/306,828
; PRIOR FILING DATE: 1999-05-07
; PRIOR APPLICATION NUMBER: US 09/227,881
; PRIOR FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 1
; LENGTH: 5300
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-244-633-1

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Query Match 99.5%; Score 5246.4; DB 43; Length 5300;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 5269; Conservative 0; Mismatches 1; Indels 2; Gaps 2;
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Db 1 ATCTTTGTTTCAGTTTACCTCAGGGCTATTATGAAATGAAATGAGATAACCAATGTGAAG 60

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Db 61 TCCTATAAAGCTATATAGCTCCATTCGGATGTATGTCTTTGGCAGGATGATAAAGATCA 120
Qy 121 GGAAGAAGGAGTATCCACGTTAGCCAAAGTGTCCAGAGGTGTCTGTCTTATTTTAGTGA 180
Db 121 GGAAGAAGGAGTATCCACGTTAGCCAAAGTGTCCAGAGGTGTCTGTCTTATTTTAGTGA 180
Qy 181 CAGATGTTGCTCTCGACAGAGCTATTTCTTCAGGAAACATCATCATCATATGTTAAATC 240
Db 181 CAGATGTTGCTCTCGACAGAGCTATTTCTTCAGGAAACATCATCATCATATGTTAAATC 240
Qy 241 CATCAACAGAGCTAAGAAACAGGATGAGTGGGCACTTGGCCCAAGGAAAAATGCCAG 300
Db 241 CATCAACAGAGCTAAGAAACAGGATGAGTGGGCACTTGGCCCAAGGAAAAATGCCAG 300
Qy 301 GAGAGCAAAATTAATGATGAAAAATAAACTTTTCCCTTTTGTGTTTAAATTTTCAGAAAAAATG 360
Db 301 GAGAGCAAAATTAATGATGAAAAATAAACTTTTCCCTTTTGTGTTTAAATTTTCAGAAAAAATG 360
Qy 361 ATGAGGCCAAATCAATGAATGAAGAAACAGCTCAGAAAAAAGATGTTTCCAAATGG 420
Db 361 ATGAGGCCAAATCAATGAATGAAGAAACAGCTCAGAAAAAAGATGTTTCCAAATGG 420
Qy 421 TAATTAAGTATTTGTTCTTTGGGAAGAGACCTCATGTGAGCTTGTATGTTTAAATTTTCAGAAAAAATG 480
Db 421 TAATTAAGTATTTGTTCTTTGGGAAGAGACCTCATGTGAGCTTGTATGTTTAAATTTTCAGAAAAAATG 480
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Db 721 ATTGACTGGGCTAAGCCTGACCTTTCAAGGAAATATGAAAAACTGAGAGCAACAAAA 780
Qy 781 GACATGTTTAAAGGCAACACAGAACATTTGAGCCCTTCAAGCAGCAGTCCCTCAGCA 840
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Qy 1021 GGATAGGTCAGAAATCATTTAGAAATCCTGTGTGCCCATCTTAACTTTTTCAGAAATGATC 1080
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Qy 4980 GTGTAAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 5039
Db 4980 GTGTAAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 5039
Qy 5040 ATAGGAATCTAATTTGGGTATGGGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5099
Db 5040 ATAGGAATCTAATTTGGGTATGGGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5099
Qy 5100 CCAACAGACTCTGGAAGGTTATTTTCTAAGAAATCTTGGTGGAGGCTGGAAGCAACCC 5159
Db 5100 CCAACAGACTCTGGAAGGTTATTTTCTAAGAAATCTTGGTGGAGGCTGGAAGCAACCC 5159
Qy 5160 CCTGTGCAAGCCCAAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 5219
Db 5160 CCTGTGCAAGCCCAAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 5219
Qy 5220 GGCTCCCCAGTATATATAACCTCTCTGAGGCTCGGGCATGAGCCAGCAAGG 5271
Db 5220 GGCTCCCCAGTATATATAACCTCTCTGAGGCTCGGGCATGAGCCAGCAAGG 5271

RESULT 10

US-10-741-339-1

; Sequence 1, Application US/10741339

; GENERAL INFORMATION:

; APPLICANT: Polansky, Jon

; TITLE OF INVENTION: METHODS TO SCREEN AND TREAT INDIVIDUALS WITH GLAUCOMA OR THE P

; FILE REFERENCE: 13587.375

; CURRENT APPLICATION NUMBER: US/10/741,339

; PRIOR FILING DATE: 2003-12-22

; PRIOR APPLICATION NUMBER: US 09/985,637

; PRIOR FILING DATE: 2001-11-05

; NUMBER OF SEQ ID NOS: 21

; SOFTWARE: PatentIn version 3.1

; SEQ ID NO 1

; LENGTH: 5300
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-741-339-1

Query Match 99.5%; Score 5246.4; DB 61; Length 5300;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 5269; Conservative 0; Mismatches 1; Indels 2; Gaps 2;

961	Qy	TTT	TAG	CAT	GGGTCCCAATTTT	TATAAGTCAGGCATACAAAGGATAACGTGTGCCAGCTCC	1020
961	Db	TTT	TAG	CAT	GGGTCCCAATTTT	TATAAGTCAGGCATACAAAGGATAACGTGTGCCAGCTCC	1020
1021	Qy	GGAT	AGGT	CAGAAATCA	TTAGAAATC	ACGTGTGTCCCACTCTAACTTTTTCAGAAATGATC	1080
1021	Db	GGAT	AGGT	CAGAAATCA	TTAGAAATC	ACGTGTGTCCCACTCTAACTTTTTCAGAAATGATC	1080
1081	Qy	TGTC	ATAG	CCCTCAC	CACAGG	CCGATGTGTGACCTTACAACCATCTACAACCCAA	1140
1081	Db	TGTC	ATAG	CCCTCAC	CACAGG	CCGATGTGTGACCTTACAACCATCTACAACCCAA	1140
1141	Qy	GTCC	CTCA	ACCAATG	TTAACTGT	CACTCAGTAGTCCCATTTACAANTGCGACCTCCCC	1200
1141	Db	GTCC	CTCA	ACCAATG	TTAACTGT	CACTCAGTAGTCCCATTTACAANTGCGACCTCCCC	1200
1201	Qy	TGTC	AGCCCAT	CCCGCT	CCACAGG	AAAGTCTCCCACTCTAGACATTTCTGCATCACGATGT	1260
1201	Db	TGTC	AGCCCAT	CCCGCT	CCACAGG	AAAGTCTCCCACTCTAGACATTTCTGCATCACGATGT	1260
1261	Qy	TACAG	CAGA	AGCTCC	GTGAGG	TGAGGGTCTGTGCTTTACACCTACTGTATGCTCTAC	1320
1261	Db	TACAG	CAGA	AGCTCC	GTGAGG	TGAGGGTCTGTGCTTTACACCTACTGTATGCTCTAC	1320
1321	Qy	ACCT	GAGCTC	ACTG	GAACCT	CTGCTCCAGGTTCAAGCAATTTCTCCTCTCTCAGCCTCC	1380
1321	Db	ACCT	GAGCTC	ACTG	GAACCT	CTGCTCCAGGTTCAAGCAATTTCTCCTCTCTCAGCCTCC	1380
1381	Qy	CGGT	AGCTGG	GA	CTACAG	CGCACGCCCGGCTAAATTTTGTATTTAGTAGTAGAGATGG	1440
1381	Db	CGGT	AGCTGG	GA	CTACAG	CGCACGCCCGGCTAAATTTTGTATTTAGTAGTAGAGATGG	1440
1441	Qy	GTTC	ACCA	TAAT	TAGCCCG	CTGGTCTTGAACCTCTGACCTCAGGTGATCCACCCACCTC	1500
1441	Db	GTTC	ACCA	TAAT	TAGCCCG	CTGGTCTTGAACCTCTGACCTCAGGTGATCCACCCACCTC	1500
1501	Qy	AGCT	CTTAA	AGTCTG	GGATAC	AGCGATGAGTACCGGCCCGGCCAAGGTCAGTGT	1560
1501	Db	AGCT	CTTAA	AGTCTG	GGATAC	AGCGATGAGTACCGGCCCGGCCAAGGTCAGTGT	1560
1561	Qy	TTAAT	AGGA	TAAC	TTGAAT	TGTTTACTAAACCAACAGGGAAACAGACAAAGCTGTGA	1620
1561	Db	TTAAT	AGGA	TAAC	TTGAAT	TGTTTACTAAACCAACAGGGAAACAGACAAAGCTGTGA	1620
1621	Qy	TAAT	TT	CAGG	ATCTTGG	ATCGGGAAATGGTGCCATGAGCTGCCTGCTAGTCCCGAGC	1680
1621	Db	TAAT	TT	CAGG	ATCTTGG	ATCGGGAAATGGTGCCATGAGCTGCCTGCTAGTCCCGAGC	1680
1681	Qy	CAC	TGGT	CTCAT	CACTTCT	CCCTCATCTCAATTTTCAGGCTAAGTTACCAATTTTATT	1740
1681	Db	CAC	TGGT	CTCAT	CACTTCT	CCCTCATCTCAATTTTCAGGCTAAGTTACCAATTTTATT	1740
1741	Qy	CAC	CA	AGCTTTT	TG	GTGTAAGCTTCAATCTGTAATTAAGATATACATAAATAG	1800
1741	Db	CAC	CA	AGCTTTT	TG	GTGTAAGCTTCAATCTGTAATTAAGATATACATAAATAG	1800
1801	Qy	TTCC	ATT	TGG	CCATCT	GTGTGTATAGGGAGGAGGCATACCCACAGAGACTCCT	1860
1801	Db	TTCC	ATT	TGG	CCATCT	GTGTGTATAGGGAGGAGGCATACCCACAGAGACTCCT	1860
1861	Qy	TGA	AG	CCCCCG	CAGAG	GTTCCTCTCCAGCTGGGGAGCCCTTCAAGCACCCGGGGTCC	1920
1861	Db	TGA	AG	CCCCCG	CAGAG	GTTCCTCTCCAGCTGGGGAGCCCTTCAAGCACCCGGGGTCC	1920
1921	Qy	TGG	GT	GTCT	GAG	AACTGCGAGCGGTGCCACTGGTTGTTTGTATACACTCTTAG	1980
1921	Db	TGG	GT	GTCT	GAG	AACTGCGAGCGGTGCCACTGGTTGTTTGTATACACTCTTAG	1980
1981	Qy	GAC	CT	GT	GTCT	TATTTCTGTGTACTCGTTTCAATTCACGAGCATTCATTGACAAAT	2040
1981	Db	GAC	CT	GT	GTCT	TATTTCTGTGTACTCGTTTCAATTCACGAGCATTCATTGACAAAT	2040
2041	Qy	TATT	GAG	TACT	TATAT	CTGCCAGACACAGAGACAAATGGTGAGCAAAAGCTCACTGC	2100

DB	2041	TATTGAGTACTTATATCTTGCAGACACACAGAGACAAATGGTGTAGCAAAAGCAGTCACTGC	2100
QY	2101	CTTACTCTTCTGGAGGTGACAGTTTCTCATGGAAAGCGTCAGAAAGAAAATTAATAGCCA	2160
DB	2101	CTTACTCTTCTGGAGGTGACAGTTTCTCATGGAAAGCGTCAGAAAGAAAATTAATAGCCA	2160
QY	2161	GCCAACTTAAACCCAGTCTCTGAAAGAAAGGAAAATAAACACCATCTTGAAGAAATTTGTCGC	2220
DB	2161	GCCAACTTAAACCCAGTCTCTGAAAGAAAGGAAAATAAACACCATCTTGAAGAAATTTGTCGC	2220
QY	2221	AGCATCCCTTAAACAGGCCACCTCCCTTAGCGCCCTGCTGCTCCTCATCTGTCGCCGAGG	2280
DB	2221	AGCATCCCTTAAACAGGCCACCTCCCTTAGCGCCCTGCTGCTCCTCATCTGTCGCCGAGG	2280
QY	2281	CCCCAAGCCCGAGTCTTCCAAAGCCTCTCTCATAGTCACAGCGCTGCAGCTGSCCT	2340
DB	2281	CCCCAAGCCCGAGTCTTCCAAAGCCTCTCTCATAGTCACAGCGCTGCAGCTGSCCT	2340
QY	2341	GCCTCGCTTCCCGTGAATCGTCTGGTGATCTGAGCTGGAGACTCTTGGCTCCAGGCT	2400
DB	2341	GCCTCGCTTCCCGTGAATCGTCTGGTGATCTGAGCTGGAGACTCTTGGCTCCAGGCT	2400
QY	2401	CCAGAAAGGAAATGGAGAGGAAACTAGTCTAAACGGAGATCTCGAGCGGACAGTGTTTC	2460
DB	2401	CCAGAAAGGAAATGGAGAGGAAACTAGTCTAAACGGAGATCTCGAGCGGACAGTGTTTC	2460
QY	2461	CTCAGAGGAAAGGGCCCTCCACGTCACAGGAGAAATCCAGAGAGTGGGAGCTGCAGGAG	2520
DB	2461	CTCAGAGGAAAGGGCCCTCCACGTCACAGGAGAAATCCAGAGAGTGGGAGCTGCAGGAG	2520
QY	2521	TGGGGACGCTGGGGCTGAGCGGGTGCTGAAAGCGAGGAAGGTGAAAAGGGCAAGGCTGAA	2580
DB	2521	TGGGGACGCTGGGGCTGAGCGGGTGCTGAAAGCGAGGAAGGTGAAAAGGGCAAGGCTGAA	2580
QY	2581	GCTGCCAGATGTTCAAGTGTGTTTCAAGGGCTGGAGTTTTCCGTTGCTTCTGTTGAGC	2640
DB	2581	GCTGCCAGATGTTCAAGTGTGTTTCAAGGGCTGGAGTTTTCCGTTGCTTCTGTTGAGC	2640
QY	2641	CTTTTATCTTTTCTCTGCTTGGAGGAGAAAGTCTATTTCAATGAAGGAGTGCAGTTTC	2700
DB	2641	CTTTTATCTTTTCTCTGCTTGGAGGAGAAAGTCTATTTCAATGAAGGAGTGCAGTTTC	2700
QY	2701	ATAAGTCAGCTGTTTAAATTTCCAGGGTGTCATGGGTTTTCTTCCACGAAGGCTTTAT	2760
DB	2701	ATAAGTCAGCTGTTTAAATTTCCAGGGTGTCATGGGTTTTCTTCCACGAAGGCTTTAT	2760
QY	2761	TTAATGGGAAATAGGAAGCGAGCTCAITTCCTTAGCGCGTTAAATTCACGAAGAAGTGAC	2820
DB	2761	TTAATGGGAAATAGGAAGCGAGCTCAITTCCTTAGCGCGTTAAATTCACGAAGAAGTGAC	2820
QY	2821	TGGAGTCTTTTCTTTCACTGCTTCTGGGCAACTACTCAGCCCTGTGTGACTTGCGTTA	2880
DB	2821	TGGAGTCTTTTCTTTCACTGCTTCTGGGCAACTACTCAGCCCTGTGTGACTTGCGTTA	2880
QY	2881	TGCAAGACGCTCGAAAAACCTTTGGAATCAGGAGACTCGGTTTTCTTCTGTGTTCTGCCATT	2940
DB	2881	TGCAAGACGCTCGAAAAACCTTTGGAATCAGGAGACTCGGTTTTCTTCTGTGTTCTGCCATT	2940
QY	2941	GGTTGGCTGTGCGACCGTGGGCAAGTGTCTCTCTTCCCTGGGCCATPAGTCTTCTCTGCT	3000
DB	2941	GGTTGGCTGTGCGACCGTGGGCAAGTGTCTCTCTTCCCTGGGCCATPAGTCTTCTCTGCT	3000
QY	3001	ATAAAGACCTTTCAGCTCTCGTGTCTGTGAAACACTTCCCTGTGATCTCTGTGAGGG	3060
DB	3001	ATAAAGACCTTTCAGCTCTCGTGTCTGTGAAACACTTCCCTGTGATCTCTGTGAGGG	3060
QY	3061	GGATGTTTGAGAGGGGAAAGGAGCGCAGAGCTGGAGCAGCTGAGCCACAGGGGAGGTGGAGG	3120
DB	3061	GGATGTTTGAGAGGGGAAAGGAGCGCAGAGCTGGAGCAGCTGAGCCACAGGGGAGGTGGAGG	3120
QY	3121	GGACAGGAAGGCGAGCGAAGACTGGGTGCTCCATPAGTCTCTCACTGATCATCGTCAGATC	3180

Db	3121	GGACGAGGACGAGCAGAGAGCTGGGTGCTCCATCAGTCCTCACTGATCAGCTCAGACTC	3180
Qy	3181	CAGGACCGAGAGCCACAATGCTTCAGGAAAGCTCAATGAACCCCAACAGCCACATTTTCCT	3240
Db	3181	CAGGACCGAGAGCCACAATGCTTCAGGAAAGCTCAATGAACCCCAACAGCCACATTTTCCT	3240
Qy	3241	TCCCTAAGCATAGACAATGGCAATTGTCCAATAACCAAAAAAGATGCAGAGACTAACTGGT	3300
Db	3241	TCCCTAAGCATAGACAATGGCAATTGTCCAATAACCAAAAAAGATGCAGAGACTAACTGGT	3300
Qy	3301	GGTAGCTTTTGCCTGGCATTTCAAAAACCTGGCCAGAGCAGCTGGAAATGCCAGAGATTG	3360
Db	3301	GGTAGCTTTTGCCTGGCATTTCAAAAACCTGGCCAGAGCAGCTGGAAATGCCAGAGATTG	3360
Qy	3361	TTAAACTTTTCCCTGACCAAGCACCACGACGCTCAGCAGTGACTGCTGAAGCAACGG	3420
Db	3361	TTAAACTTTTCCCTGACCAAGCACCACGACGCTCAGCAGTGACTGCTGAAGCAACGG	3420
Qy	3421	AGTGACTGACGGCAGGGGAGGAGAAAAAGAGAGGGATAGTGATGAGCAAGAAAG	3480
Db	3421	AGTGACTGACGGCAGGGGAGGAGAAAAAGAGAGGGATAGTGATGAGCAAGAAAG	3480
Qy	3481	ACGATTTCAATCAAGGCGCTGGGAATGACCAAGGATTTATAGTCCACGCTGATCCTGG	3540
Db	3481	ACGATTTCAATCAAGGCGCTGGGAATGACCAAGGATTTATAGTCCACGCTGATCCTGG	3540
Qy	3541	GTTCTAGAGCGCAGGGCTATATTTGTGGGGGAAAAAATCAGTTCAAGGGAAGTCGGAGA	3600
Db	3541	GTTCTAGAGCGCAGGGCTATATTTGTGGGGGAAAAAATCAGTTCAAGGGAAGTCGGAGA	3600
Qy	3601	CCTGATTTCTAATACTATATTTTTCCTTTTCAAGCTGAGTAAATCTTGAGCAAGTCAAG	3660
Db	3601	CCTGATTTCTAATACTATATTTTTCCTTTTCAAGCTGAGTAAATCTTGAGCAAGTCAAG	3660
Qy	3661	GTAGTAATCAGGCTGTAAAGTTACTTAGTTTCTCCTTATTAGGAATCTTTTTCCTCTGT	3720
Db	3661	GTAGTAATCAGGCTGTAAAGTTACTTAGTTTCTCCTTATTAGGAATCTTTTTCCTCTGT	3720
Qy	3721	GGAGTTAGCAGCAAGGGGCAATCCCGTTTCTTTTAAACAGGAAGAAACATTCCTAAGAG	3780
Db	3721	GGAGTTAGCAGCAAGGGGCAATCCCGTTTCTTTTAAACAGGAAGAAACATTCCTAAGAG	3780
Qy	3781	TAAAGCCAAACAGATCAAGCTTAGTCTTGCTGACTATATGATGGTTTTTTTGA AAAAT	3840
Db	3781	TAAAGCCAAACAGATCAAGCTTAGTCTTGCTGACTATATGATGGTTTTTTTGA AAAAT	3840
Qy	3841	CATTTGAGCGATGTTACTATCTGATTCAGAAAATGAGACTAGTACCCCTTGCTGAGCTG	3900
Db	3841	CATTTGAGCGATGTTACTATCTGATTCAGAAAATGAGACTAGTACCCCTTGCTGAGCTG	3900
Qy	3901	TAAACAAACCCAGTTGTAAATGCTCAGTTTCAGGCTTAACTGCAGAGACCAATCAAA-	3959
Db	3901	TAAACAAACCCAGTTGTAAATGCTCAGTTTCAGGCTTAACTGCAGAGACCAATCAAA-	3959
Qy	3960	AAGAAATAGAAATCTTTAGAGCAAACTGTGTTTCTCCACATCTGGAGGTGAGTCTGCCAGG	4019
Db	3961	AAGAAATAGAAATCTTTAGAGCAAACTGTGTTTCTCCAC-TCGGAGGTGAGTCTGCCAGG	4019
Qy	4020	CAGTTTGGAAATATTTACTTTCACAAGTATGACACTGTTGTTGGTATTAACAACATAAAG	4079
Db	4020	CAGTTTGGAAATATTTACTTTCACAAGTATGACACTGTTGTTGGTATTAACAACATAAAG	4079
Qy	4080	TTGCTCAAAGGCAATCATTTATTTCAAGTCGCTTAAAGTTACTTCTCAGAGTTTGTGTATA	4139
Db	4080	TTGCTCAAAGGCAATCATTTATTTCAAGTCGCTTAAAGTTACTTCTCAGAGTTTGTGTATA	4139
Qy	4140	TTTATTTGGCTATTTGCCATTTGCTTTTTTGTTTTTTCTTTGGGTTTTTAAATGAAGCA	4199
Db	4140	TTTATTTGGCTATTTGCCATTTGCTTTTTTGTTTTTTCTTTGGGTTTTTAAATGAAGCA	4199
Qy	4200	GGGATTTAAACCTACAGTCCAGAAAGCCTGTGAAATTTGAAATGAGGAAAAAATTAATTT	4259
Db	4200	GGGATTTAAACCTACAGTCCAGAAAGCCTGTGAAATTTGAAATGAGGAAAAAATTAATTT	4259


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4260 TTGTTTTTACCACCTTCTAACTAAATTTAACTATTTTATTCCTATGCGAATAGAGCCATAA 4319
4260 TTGTTTTTACCACCTTCTAACTAAATTTAACTATTTTATTCCTATGCGAATAGAGCCATAA 4319
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4440 ACTTTGAAATTTAGACCTCTCTGCTGGATCTTTGTTTAAACATATTAATAAAAAATGTTTAA 4499
4440 ACTTTGAAATTTAGACCTCTCTGCTGGATCTTTGTTTAAACATATTAATAAAAAATGTTTAA 4499
4500 AATTTTGATATTTTGATATATCATATTTCAATATTCATTTGTTTCTTTGTAATCTATATTT 4559
4500 AATTTTGATATTTTGATATATCATATTTCAATATTCATTTGTTTCTTTGTAATCTATATTT 4559
4560 TATATATTTGAAACATCTTTCTGAGAAGAGTTCCCGAGATTTCCACCAATGAGGTTCTTG 4619
4560 TATATATTTGAAACATCTTTCTGAGAAGAGTTCCCGAGATTTCCACCAATGAGGTTCTTG 4619
4620 GCATGCACACACACAGAGTAAGAACTGATTTAGAGGCTAACATTTGATGATGTCCTGAG 4679
4620 GCATGCACACACACAGAGTAAGAACTGATTTAGAGGCTAACATTTGATGATGTCCTGAG 4679
4680 ATGCAAGACTGAAATTTAGAAAGTTCTCCCAAGATACACAGTTGTTTAAAGCTAGGGCT 4739
4680 ATGCAAGACTGAAATTTAGAAAGTTCTCCCAAGATACACAGTTGTTTAAAGCTAGGGCT 4739
4740 GAGGGGGGAAATCTCGCGCTTCTATAGGAATGCTCTCCCTGGAGCTGCTGAGGCTGCTGT 4799
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4800 CCTTGTTGTTCTGGCTGGCTGTTATTTTCTCTGCTGCTGCTACGTTCTTAAAGGACTTTGT 4859
4800 CCTTGTTGTTCTGGCTGGCTGTTATTTTCTCTGCTGCTGCTACGTTCTTAAAGGACTTTGT 4859
4860 TGATCTCAGATTCTTAGCATAGTGTGCTGCGACAGTGCAGGTTCTCAATGAGTTGCGAG 4919
4860 TGATCTCAGATTCTTAGCATAGTGTGCTGCGACAGTGCAGGTTCTCAATGAGTTGCGAG 4919
4920 GTCAATGGAATATAAATAAGAAATATATCTTTGTTGAATCAGACACACAGTAGTCTTG 4979
4920 GTCAATGGAATATAAATAAGAAATATATCTTTGTTGAATCAGACACACAGTAGTCTTG 4979
4980 GTGTAAGTGTGTGACGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 5039
4980 GTGTAAGTGTGTGACGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 5039
5040 ATAGGAACATAATTATGGGGTATGGGTGCATAAATTTGGATGTTCTTTTAAAGAACT 5099
5040 ATAGGAACATAATTATGGGGTATGGGTGCATAAATTTGGATGTTCTTTTAAAGAACT 5099
5100 CCAACAGACTCTCGGAAGTTATTTTCTAAGAACTCTGCTGCGAGCTGAGGCAACCC 5159
5100 CCAACAGACTCTCGGAAGTTATTTTCTAAGAACTCTGCTGCGAGCTGAGGCAACCC 5159
5160 CCTGTGTCACAGCCCCACCGAGCTCAGTGGCCACCTCTGTCTTCCCCCATGAGGGCT 5219
5160 CCTGTGTCACAGCCCCACCGAGCTCAGTGGCCACCTCTGTCTTCCCCCATGAGGGCT 5219
5220 GGCTCCCCAGTATATATAAACCTCTCTGGAGCTCGGCGATGAGCCAGCAGG 5271
5220 GGCTCCCCAGTATATATAAACCTCTCTGGAGCTCGGCGATGAGCCAGCAGG 5271
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RESULT 11

US-10-087-192-1228

; Sequence 1228, Application US/10087192

; GENERAL INFORMATION:

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; APPLICANT: Morris, David W.
; APPLICANT: Engelhard, Eric K.
; TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
; TITLE OF INVENTION: CANCER
; FILE REFERENCE: 529452000122
; CURRENT APPLICATION NUMBER: US/10/087,192
; CURRENT FILING DATE: 2002-03-01
; PRIOR APPLICATION NUMBER: US 09/747,377
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 09/798,586
; PRIOR FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 2059
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1228
; LENGTH: 37252
; TYPE: DNA
; ORGANISM: Homo sapiens
; US-10-087-192-1228

Query Match 99.3%; Score 5232.4; DB 40; Length 37252;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 5267; Conservative 0; Mismatches 1; Indels 4; Gaps 3;

QY 1 ATCTTTGTTCAAGTTTACCTCAGGGCTATTATGAATGAAATGAGATAACCAATGTGAAAG 60
DB 4731 ATCTTTGTTCAAGTTTACCTCAGGGCTATTATGAATGAAATGAGATAACCAATGTGAAAG 4790
QY 61 TCCTATAAATCTGTATAGCTCCATTCGGATGTATGCTTTTGGCAGGATGATAAGAAATCA 120
DB 4791 TCCTATAAATCTGTATAGCTCCATTCGGATGTATGCTTTTGGCAGGATGATAAGAAATCA 4850
QY 121 GGAAGAAGAGATATCCAGTTAGCCAGTGTCCAGGCTGTGTCTGCTCTTATTTTAGTGA 180
DB 4851 GGAAGAAGAGATATCCAGTTAGCCAGTGTCCAGGCTGTGTCTGCTCTTATTTTAGTGA 4910
QY 181 CAGATGTTGCTCTCAGCAGAGAGCTATTCTTCAGGAAACATCATCATTAATGTTAAATC 240
DB 4911 CAGATGTTGCTCTCAGCAGAGAGCTATTCTTCAGGAAACATCATCATTAATGTTAAATC 4970
QY 241 CATCAAAACAGGAGCTTAAGAAAACAGGAATGAGATGGGCACTTCCCAAGAAAAATGCCAG 300
DB 4971 CATCAAAACAGGAGCTTAAGAAAACAGGAATGAGATGGGCACTTCCCAAGAAAAATGCCAG 5030
QY 301 GAGACAAATAATGATGAAAAATAAACTTTTCCCTTTGTTTAAATTTTTCAGGAAAAATG 360
DB 5031 GAGACAAATAATGATGAAAAATAAACTTTTCCCTTTGTTTAAATTTTTCAGGAAAAATG 5090
QY 361 ATGAGGACCAAAATCAATGAATAAGAAAAACAGCTCAGAAAAAAGATGTTTCCAAATTTGG 420
DB 5091 ATGAGGACCAAAATCAATGAATAAGAAAAACAGCTCAGAAAAAAGATGTTTCCAAATTTGG 5150
QY 421 TAATTAAGTATTTGTTCTTGGGAAAGAGACCTCCATGTGAGCTTGATGGGAAAAATGGGAA 480
DB 5151 TAATTAAGTATTTGTTCTTGGGAAAGAGACCTCCATGTGAGCTTGATGGGAAAAATGGGAA 5210
QY 481 AAACGTCAAAACAGTATCTGATCAGATCCCAAAGTGGATTAATTTTAAACACAGAT 540
DB 5211 AAACGTCAAAACAGTATCTGATCAGATCCCAAAGTGGATTAATTTTAAACACAGAT 5270
QY 541 GGCATCATCTCTGGGAGGCAAGTTCCAGGAAGTCTATGTTAGCAAAAGGACATTAACATAAC 600
DB 5271 GGCATCATCTCTGGGAGGCAAGTTCCAGGAAGTCTATGTTAGCAAAAGGACATTAACATAAC 5330
QY 601 AGCAAAATCAAAATTTCCGCAAAATGCAGGAGGAAAAATGGGAGCTGGGAAAAAGCTTTTCAAC 660
DB 5331 AGCAAAATCAAAATTTCCGCAAAATGCAGGAGGAAAAATGGGAGCTGGGAAAAAGCTTTTCAAC 5390
QY 661 AGTGATTAGCGAGTTGA CCAATGTTTCGCAACACTCCCGCTCTATATACCAGGGAACACAAAA 720
DB 5391 AGTGATTAGCGAGTTGA CCAATGTTTCGCAACACTCCCGCTCTATATACCAGGGAACACAAAA 5450
QY 721 ATTGACTGGGCTAAGCCTGAGCTTTTCAAGGGAAATATGAAAACTTGAGAGCAAAACAAAA 780
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Db 5451 ATTGACTGGGCTAAGCCCTGGACTTTCAAGGGAATATGAAGAACTGAGAGCAAAACAAA 5510
Qy 781 GACATGGTTAAAGCAACCGAACAATTTGTGAGCCCTCAAAGCAGCAGTGCCCTCAGCA 840
Db 5511 GACATGGTTAAAGCAACCGAACAATTTGTGAGCCCTCAAAGCAGCAGTGCCCTCAGCA 5570
Qy 841 GGGACCCCTGAGGCAATTTGCCCTTTAGGAAGGCCAGTTTCTTAAGGAATCTTAAGAACTC 900
Db 5571 GGGACCCCTGAGGCAATTTGCCCTTTAGGAAGGCCAGTTTCTTAAGGAATCTTAAGAACTC 5630
Qy 901 TTGAAGATCATGAATTTTAAACCAATTTAAAGTATAAAACAAATATGCGATGCAATAATCAG 960
Db 5631 TTGAAGATCATGAATTTTAAACCAATTTTAAAGTATAAAACAAATATGCGATGCAATAATCAG 5690
Qy 961 TTTAGACATGGGTCCCAATTTTATAAGTCAAGGCATACAAGGATAAAGTGCCAGCTCC 1020
Db 5691 TTTAGACATGGGTCCCAATTTTATAAGTCAAGGCATACAAGGATAAAGTGCCAGCTCC 5750
Qy 1021 GGATAGGTCAAGAAATCATTAAGAAATCACTGTGTCCCAATCTTAACTTTTTCAGAAATGATC 1080
Db 5751 GGATAGGTCAAGAAATCATTAAGAAATCACTGTGTCCCAATCTTAACTTTTTCAGAAATGATC 5810
Qy 1081 TGTCAATAGCCCTCACACAGCCCGGATGTGTCTGACCTACACCAATCTTACACCCAA 1140
Db 5811 TGTCAATAGCCCTCACACAGCCCGGATGTGTCTGACCTACACCAATCTTACACCCAA 5870
Qy 1141 GTGCTCAACCAATTTTAACTGTGTATCTCAGTAGTCCCAATTAAGATGCACTCC 1200
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Qy 1201 TGTGAGCCCAATCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCAATCAGATGT 1260
Db 5931 TGTGAGCCCAATCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCAATCAGATGT 5990
Qy 1261 TACAGCCAGAACTCCGCTGAGGGTGAGGGTCTGTCTTACCACTACCTGTATCTCTAC 1320
Db 5991 TACAGCCAGAACTCCGCTGAGGGTGAGGGTCTGTCTTACCACTACCTGTATCTCTAC 6050
Qy 1321 ACCTGAGTCACTGCAACCTTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 1380
Db 6051 ACCTGAGTCACTGCAACCTTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 6110
Qy 1381 CGCTGAGTGGGACTACAGGGCGCACCGCGCTAAATTTTGTATTTGTAGTAGAGTGG 1440
Db 6111 CGCTGAGTGGGACTACAGGGCGCACCGCGCTAAATTTTGTATTTGTAGTAGAGTGG 6170
Qy 1441 GTTTCACCAATATTAGCCCGGCTGTCTTGAACCTCTGACCTCAGGTGATCCACCACTC 1500
Db 6171 GTTTCACCAATATTAGCCCGGCTGTCTTGAACCTCTGACCTCAGGTGATCCACCACTC 6230
Qy 1501 AGCTCTCTAAGTCTGGGATTAAGGATGAGTCAAGGCGCCCGGCAAGGGTCAAGTGT 1560
Db 6231 AGCTCTCTAAGTCTGGGATTAAGGATGAGTCAAGGCGCCCGGCAAGGGTCAAGTGT 6290
Qy 1561 TTAATAAGGAATTAATTAAGTGTAACTAAACCAAGGGAACACACAAAGAGCTTGA 1620
Db 6291 TTAATAAGGAATTAATTAAGTGTAACTAAACCAAGGGAACACACAAAGAGCTTGA 6350
Qy 1621 TAAATTCAGGAATCTTTGGGATGGGAATGGTGCATGAGCTGCTGCTAGTCCCGAC 1680
Db 6351 TAAATTCAGGAATCTTTGGGATGGGAATGGTGCATGAGCTGCTGCTAGTCCCGAC 6410
Qy 1681 CACTGGTCTCATCATCTTCTCCCTCATCTCTCATTTTCAAGGCTAAGTTACCAATTTAT 1740
Db 6411 CACTGGTCTCATCATCTTCTCCCTCATCTCTCATTTTCAAGGCTAAGTTACCAATTTAT 6470
Qy 1741 CACATGCTTTTGTGGTAAGCTCCACATGTTTACTGAAATTAAGAGTATACATAAATAG 1800
Db 6471 CACATGCTTTTGTGGTAAGCTCCACATGTTTACTGAAATTAAGAGTATACATAAATAG 6530
Qy 1801 TTCCATTTGGGGCCATCTGTGTGTATAGGGAGGAGGAGGATACCCAGAGACTCCT 1860

Db 6531 TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGGAGGAGGATACCCAGAGACTCCT 6590
Qy 1861 TGAAGCCCCCGGAGAGAGTTTCTCTCAGCTGGGGAGCCCTCAAGACACCCGGGGTCC 1920
Db 6591 TGAAGCCCCCGGAGAGAGTTTCTCTCAGCTGGGGAGCCCTCAAGACACCCGGGGTCC 6650
Qy 1921 TGGGTGTCTTGAGCAACCTGCGAGCCCGTGCACCTGTTTGTGTTTGTATCACTCTCTAGG 1980
Db 6651 TGGGTGTCTTGAGCAACCTGCGAGCCCGTGCACCTGTTTGTGTTTGTATCACTCTCTAGG 6710
Qy 1981 GACCTGTGTCTTCTATTTCTGTGTGACTCGTTCATTCATCAGGCAATTCATTCAGCAAT 2040
Db 6711 GACCTGTGTCTTCTATTTCTGTGTGACTCGTTCATTCATCAGGCAATTCATTCAGCAAT 6770
Qy 2041 TATTGAGTACTTATATCTGCGAGACACAGAGACAAATGGTGAAGCAAGCAGTCACTGC 2100
Db 6771 TATTGAGTACTTATATCTGCGAGACACAGAGACAAATGGTGAAGCAAGCAGTCACTGC 6830
Qy 2101 CCTACCTTCTGTGAGGTGACAGTTTCTCATGGAAGAGCTGCAGAAAGAAATTAATAGCA 2160
Db 6831 CCTACCTTCTGTGAGGTGACAGTTTCTCATGGAAGAGCTGCAGAAAGAAATTAATAGCA 6890
Qy 2161 GCCAATCTAAACCCAGTGTGAAAGAAAGAAATTAACACCATCTTGAAGAAATTTGTGCG 2220
Db 6891 GCCAATCTAAACCCAGTGTGAAAGAAAGAAATTAACACCATCTTGAAGAAATTTGTGCG 6950
Qy 2221 AGCATCCCTTAAACAGGCCACCTCCCTAGGCCCTCTGCTGCTCATCTGTCGCCGAGG 2280
Db 6951 AGCATCCCTTAAACAGGCCACCTCCCTAGGCCCTCTGCTGCTCATCTGTCGCCGAGG 7010
Qy 2281 CCCCAAGCCCGAGTCTTCCAGCCCTCTCTCATCAGTCAAGCCCTCAGCTGCGCT 2340
Db 7011 CCCCAAGCCCGAGTCTTCCAGCCCTCTCTCATCAGTCAAGCCCTCAGCTGCGCT 7070
Qy 2341 GCCTCGCTTCCCGTGAATCTGCTGTGTGCACTCTGAGCTGAGAGTCTCTTGGCTCAGGCT 2400
Db 7071 GCCTCGCTTCCCGTGAATCTGCTGTGTGCACTCTGAGCTGAGAGTCTCTTGGCTCAGGCT 7130
Qy 2401 CCAGAAAGGAATGGAGAGGAACTAGTCTAAACGGAGAACTCTGGAGGGGACAGTGTTC 2460
Db 7131 CCAGAAAGGAATGGAGAGGAACTAGTCTAAACGGAGAACTCTGGAGGGGACAGTGTTC 7190
Qy 2461 CTCAAGAGGAAAGGGCCCTCCAGTCCAGGAGAAATCCAGAGGTGGGAGTCCAGGAG 2520
Db 7191 CTCAAGAGGAAAGGGCCCTCCAGTCCAGGAGAAATCCAGAGGTGGGAGTCCAGGAG 7250
Qy 2521 TGGGAGCGCTGGGGCTGAGCGGTGCTGAAAGGACAGGAGTGAAGGGCAAGGCTGAA 2580
Db 7251 TGGGAGCGCTGGGGCTGAGCGGTGCTGAAAGGACAGGAGTGAAGGGCAAGGCTGAA 7310
Qy 2581 GCTGCCAGATGTTTCACTGTGTGTTTCAAGGGGCTGGGAGTTCCTGTTCTTCTGTGAGC 2640
Db 7311 GCTGCCAGATGTTTCACTGTGTGTTTCAAGGGGCTGGGAGTTCCTGTTCTTCTGTGAGC 7370
Qy 2641 CTTTTTATCTTTTCTCTGCTTGGAGAGAGAGTCTATTTCATGAAGGATCAGTTC 2700
Db 7371 CTTTTTATCTTTTCTCTGCTTGGAGAGAGAGTCTATTTCATGAAGGATCAGTTC 7430
Qy 2701 ATAAAGTCACTGTTTAAATTTCCAGGGTGTGCAATGGGTTTTCCTTCAAGAGGCTTTAT 2760
Db 7431 ATAAAGTCACTGTTTAAATTTCCAGGGTGTGCAATGGGTTTTCCTTCAAGAGGCTTTAT 7490
Qy 2761 TTAATGGGAATATAGGAAGCAGCTCATTTCTAGCCGTTAATTCACGGAAGAGTGAC 2820
Db 7491 TTAATGGGAATATAGGAAGCAGCTCATTTCTAGCCGTTAATTCACGGAAGAGTGAC 7550
Qy 2821 TGGAGTCTTTCTTTTCTCTGCTTGGGCAACTACTCAGCCCTGTGCTGAGCTTGCCTTA 2880
Db 7551 TGGAGTCTTTCTTTTCTCTGCTTGGGCAACTACTCAGCCCTGTGCTGAGCTTGCCTTA 7610
Qy 2881 TGCAGAGCGGTGCAAAACCTTGGAAATCAGGAGCTCGGTTTCTTTCTGTTCTGCAAT 2940
Db 7611 TGCAGAGCGGTGCAAAACCTTGGAAATCAGGAGCTCGGTTTCTTTCTGTTCTGCAAT 7670

QY 1 ATCTTTGTTTACCTCAGGCTATTATGAAATCAATGAGATAACCAATGTGAAG 60
DB 1 ATCTTTGTTTACCTCAGGCTATTATGAAATCAATGAGATAACCAATGTGAAG 60
QY 61 TCCTATAAATGTATAGCTCCATTCGGATGTATGTCTTTGGCAGAGATGAATCA 120
DB 61 TCCTATAAATGTATAGCTCCATTCGGATGTATGTCTTTGGCAGAGATGAATCA 120
QY 121 GGAAGAGAGATGATCAGCTTACCAAGTGTCCAGGCTGTCTCTTTATTTAGTGA 180
DB 121 GGAAGAGAGATGATCAGCTTACCAAGTGTCCAGGCTGTCTCTTTATTTAGTGA 180
QY 181 CAGATGTGTCTCTGACAGAGCTATTCTTCAGGAAACATCAATCAATATGTTAAATC 240
DB 181 CAGATGTGTCTCTGACAGAGCTATTCTTCAGGAAACATCAATCAATATGTTAAATC 240
QY 241 CATCAACAGGAGCTAAGAAACAGGATGAGATGGGCACTTCCCAAGGAAATGCCAG 300
DB 241 CATCAACAGGAGCTAAGAAACAGGATGAGATGGGCACTTCCCAAGGAAATGCCAG 300
QY 301 GAGAGCAAAATAATGATGAAATAAATCTTTTCCCTTTGTTTAAATTTAGGAAAAATG 360
DB 301 GAGAGCAAAATAATGATGAAATAAATCTTTTCCCTTTGTTTAAATTTAGGAAAAATG 360
QY 361 ATGAGGACCAAAATCAATGAATGAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
DB 361 ATGAGGACCAAAATCAATGAATGAAGGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
QY 421 TAATTAAGTATTTGTTCTTGGGAGAGACCTCCATGTGAGCTTGTGGGAAATGGGA 480
DB 421 TAATTAAGTATTTGTTCTTGGGAGAGACCTCCATGTGAGCTTGTGGGAAATGGGA 480
QY 481 AAACGTCAAAAGCATGATCTGATCAGATCCCAAGTGGATTAATTTTAAACACCAT 540
DB 481 AAACGTCAAAAGCATGATCTGATCAGATCCCAAGTGGATTAATTTTAAACACCAT 540
QY 541 GGATCACTCTGCGGAGGCAAGTTTCAGGAAGGTCATGTTAGCAAAAGGACATAAATAC 600
DB 541 GGATCACTCTGCGGAGGCAAGTTTCAGGAAGGTCATGTTAGCAAAAGGACATAAATAC 600
QY 601 AGCAAAATCAAAATTCGCAATGTCAGGAGGAAATGGGACCTGGGAAGCTTTTCAATAC 660
DB 601 AGCAAAATCAAAATTCGCAATGTCAGGAGGAAATGGGACCTGGGAAGCTTTTCAATAC 660
QY 661 AGTGAATGAGGCTTACCATGTTTCGCAACACCTCCCGTCTATACAGGAGGAAACAAA 720
DB 661 AGTGAATGAGGCTTACCATGTTTCGCAACACCTCCCGTCTATACAGGAGGAAACAAA 720
QY 721 ATTGACTGGCTAAGCTGGAATTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA 780
DB 721 ATTGACTGGCTAAGCTGGAATTTCAAGGAAATATGAAAACTGAGAGCAAAACAAA 780
QY 781 GACATGTTAAAGGCAACAGACATGTTGAGCCTTCAAGCAGAGCTGCCCTCAGCA 840
DB 781 GACATGTTAAAGGCAACAGACATGTTGAGCCTTCAAGCAGAGCTGCCCTCAGCA 840
QY 841 GGGACCTGAGGCAATTTGCTTTAGGAGGCGAGTTTCTTAAGGAATCTTAAGAAATC 900
DB 841 GGGACCTGAGGCAATTTGCTTTAGGAGGCGAGTTTCTTAAGGAATCTTAAGAAATC 900
QY 901 TTGAAAGATCATGAATTTTAAACATTTTAAAGTAAACAAATATGCGATGCAATACAG 960
DB 901 TTGAAAGATCATGAATTTTAAACATTTTAAAGTAAACAAATATGCGATGCAATACAG 960
QY 961 TTTAGACATGGGTCCTTAAATGATCAGGATGAGGATGAGGATGAGGATGAGGATGAGG 1020
DB 961 TTTAGACATGGGTCCTTAAATGATCAGGATGAGGATGAGGATGAGGATGAGGATGAGG 1020
QY 1021 GGTAGGTGAGAAATCATGAGAAATCACTGTGTCCCATCTTAATCTTTTTCAGAAATGATC 1080
DB 1021 GGTAGGTGAGAAATCATGAGAAATCACTGTGTCCCATCTTAATCTTTTTCAGAAATGATC 1080

QY 1081 TGTCTAGCCCTCACACAGGCCCGATGTGTCTGACCTACAAACACATCTTACAAACCAA 1140
DB 1081 TGTCTAGCCCTCACACAGGCCCGATGTGTCTGACCTACAAACACATCTTACAAACCAA 1140
QY 1141 GTGCTCTCAACCAATGTTAAAGTGTCTCATCTCAGTAGGTCCCATTTACAAATGCCACCTCCC 1200
DB 1141 GTGCTCTCAACCAATGTTAAAGTGTCTCATCTCAGTAGGTCCCATTTACAAATGCCACCTCCC 1200
QY 1201 TGTGAGGCCATTCGCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGATCAGATGT 1260
DB 1201 TGTGAGGCCATTCGCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGATCAGATGT 1260
QY 1261 TACAGCCAGAGCTCCGTGAGGCTGAGGCTGTGTCTTACACCTACCTGTATGCTCTAC 1320
DB 1261 TACAGCCAGAGCTCCGTGAGGCTGAGGCTGTGTCTTACACCTACCTGTATGCTCTAC 1320
QY 1321 ACTGAGCTCACTGCAACCTCTGCGCTCCAGGTTCAAGCAATTTCTCTCTCAGCCTCC 1380
DB 1321 ACTGAGCTCACTGCAACCTCTGCGCTCCAGGTTCAAGCAATTTCTCTCTCAGCCTCC 1380
QY 1381 CGCGTAGCTGGGACTACAGGCGCACGCCCGCTAAATTTTGTATTTGTATGATGAGATGG 1440
DB 1381 CGCGTAGCTGGGACTACAGGCGCACGCCCGCTAAATTTTGTATTTGTATGATGAGATGG 1440
QY 1441 GTTTCACCATATTTAGCCGCTGTCTTGAACCTCTGAGCTCAGGTGATCCACCCACCTC 1500
DB 1441 GTTTCACCATATTTAGCCGCTGTCTTGAACCTCTGAGCTCAGGTGATCCACCCACCTC 1500
QY 1501 AGCCTCTTAAAGTGTGGGATTTACAGGCAATGATGATCAACCGCCCGCCCAAGGCTCAGTGT 1560
DB 1501 AGCCTCTTAAAGTGTGGGATTTACAGGCAATGATGATCAACCGCCCGCCCAAGGCTCAGTGT 1560
QY 1561 TTAATAAGGAATAAATTTGAATGTTTAAACCAACAGGAAACAGACAAAGAGCTGTGA 1620
DB 1561 TTAATAAGGAATAAATTTGAATGTTTAAACCAACAGGAAACAGACAAAGAGCTGTGA 1620
QY 1621 TAAATTCAGGATTTCTTGGGATGGGAATGGTCCATGAGCTGCTGCTAGTCCAGAC 1680
DB 1621 TAAATTCAGGATTTCTTGGGATGGGAATGGTCCATGAGCTGCTGCTAGTCCAGAC 1680
QY 1681 CACTGTCTCTCATCTCTTCTCCCTCATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 1740
DB 1681 CACTGTCTCTCATCTCTTCT 1740
QY 1741 CACATGCTTTTGTGTGAAGCTTCCACATGCTTACTGTAATAAGATATACATAAAGTAG 1800
DB 1741 CACATGCTTTTGTGTGAAGCTTCCACATGCTTACTGTAATAAGATATACATAAAGTAG 1800
QY 1801 TTTCCATTTGGGCGCATCTGTGTGTGTATAGGGAGGAGGCGCATACCCAGAGACTCCT 1860
DB 1801 TTTCCATTTGGGCGCATCTGTGTGTGTATAGGGAGGAGGCGCATACCCAGAGACTCCT 1860
QY 1861 TGAAGCCCGCGGAGAGGTTTCTCTCTCAGCTGGGGAGCCCTGCAAGCAACCCGGGTCC 1920
DB 1861 TGAAGCCCGCGGAGAGGTTTCTCTCTCAGCTGGGGAGCCCTGCAAGCAACCCGGGTCC 1920
QY 1921 TGGGTGCTCTGAGCAACCTGCGGCGGCTGCGACTGCTGTTGTTTGTATCAGTCTCTAGG 1980
DB 1921 TGGGTGCTCTGAGCAACCTGCGGCGGCTGCGACTGCTGTTGTTTGTATCAGTCTCTAGG 1980
QY 1981 GACCTGTCTCTTCTATTTCTGTGTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 2040
DB 1981 GACCTGTCTCTTCTATTTCTGTGTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 2040
QY 2041 TATTTAGTACTTATATCTGCCAGACACAGAGCAAAATGGTGCAGCAAAAGCAGTCTGC 2100
DB 2041 TATTTAGTACTTATATCTGCCAGACACAGAGCAAAATGGTGCAGCAAAAGCAGTCTGC 2100
QY 2101 CCTACCTTCTGTGAGGTGACGTTTCTCATGGAAGCGTGCAGAAAGAAATTAATAGCCA 2160
DB 2101 CCTACCTTCTGTGAGGTGACGTTTCTCATGGAAGCGTGCAGAAAGAAATTAATAGCCA 2160
QY 2161 GCCAACTTAAACCCAGTGTGAAAGAAAGAAATAAACCACTCTTTGAGAAATTTGTGCGC 2220

[illegible]

3241	DB	TCCCTAAGCATAGACAATGGGCATTTGGCCAAATAACAAAGAAATGCAGAGACTAATCTGGT	3300
3301	QY	GGTAGCTTTTGCCTGGCAATTCAAAACTGGCCAGACAGCAAGTGGAAAAATGCCAGAGATTG	3360
3301	DB	GGTAGCTTTTGCCTGGCAATTCAAAACTGGCCAGACAGCAAGTGGAAAAATGCCAGAGATTG	3360
3361	QY	TTAAACTTTTCCCTCTGACGACGACCCACGCGAGCTCAGCAGTGACTGTCTGACAGCACGG	3420
3361	DB	TTAAACTTTTCCCTCTGACGACGACCCACGCGAGCTCAGCAGTGACTGTCTGACAGCACGG	3420
3421	QY	AGTGACCTGACGCGCAGGGGAGGAGAAAAAGAGAGGGATAGTGTATGAGCAAGAAAG	3480
3421	DB	AGTGACCTGACGCGCAGGGGAGGAGAAAAAGAGAGGGATAGTGTATGAGCAAGAAAG	3480
3481	QY	ACAGATTCAATTCAGGGCAGTGGGAATTGACCAACAGGGATTATAGTCCACGTGATCTCTGG	3540
3481	DB	ACAGATTCAATTCAGGGCAGTGGGAATTGACCAACAGGGATTATAGTCCACGTGATCTCTGG	3540
3541	QY	GTTCTAGAGGCAAGGGCTATATTTGTGGGGGAAAAAATCAGTTCAAGGGGAAGTCGGGAGA	3600
3541	DB	GTTCTAGAGGCAAGGGCTATATTTGTGGGGGAAAAAATCAGTTCAAGGGGAAGTCGGGAGA	3600
3601	QY	CCTGATTTCTAATACTACTATATTTTCCCTTTTACAAGCTGAGTAATCTGAGCAAGTCAACAG	3660
3601	DB	CCTGATTTCTAATACTACTATATTTTCCCTTTTACAAGCTGAGTAATCTGAGCAAGTCAACAG	3660
3661	QY	GTAGTAACTGAGGCTGTAAAGATTACTTAGTTTCTCCTTATTAGGAACTCTTTTCTCTCTGT	3720
3661	DB	GTAGTAACTGAGGCTGTAAAGATTACTTAGTTTCTCCTTATTAGGAACTCTTTTCTCTCTGT	3720
3721	QY	GGAGTTAGCAGCAACAGGGCAATCCCGTTTCTTTTAAACAGGAAGAAAAATTCCTCAAGAG	3780
3721	DB	GGAGTTAGCAGCAACAGGGCAATCCCGTTTCTTTTAAACAGGAAGAAAAATTCCTCAAGAG	3780
3781	QY	TAAAGCCAAACAGATTCACGCTTAGTCTTGCTGACTATATGATGGTTTTTTTGAANAAT	3840
3781	DB	TAAAGCCAAACAGATTCACGCTTAGTCTTGCTGACTATATGATGGTTTTTTTGAANAAT	3840
3841	QY	CATTTGAGCGATGTTTACTATCTGATTCAGAAATCAGACTAGTACCCCTTGGTCAGCTG	3900
3841	DB	CATTTGAGCGATGTTTACTATCTGATTCAGAAATCAGACTAGTACCCCTTGGTCAGCTG	3900
3901	QY	TAAACAAACCCAGTTGTAAATGTCTCAAGTTTCAGGCTTAACTGCAGAACCAATCAAA-	3959
3901	DB	TAAACAAACCCAGTTGTAAATGTCTCAAGTTTCAGGCTTAACTGCAGAACCAATCAAA	3960
3960	QY	AAGAAATAGAAATCTTTAGAGCAAACTGTGTTTCTCACAATCTGGAGGTGAGTCTGCCAGGG	4019
3961	DB	AAGAAATAGAAATCTTTAGAGCAAACTGTGTTTCTCACA-TCCTGGAGGTGAGTCTGCCAGGG	4019
4020	QY	CAGTTTGGAAATATTTACTTTCACAAGTATTGACACTGTTGTGGTATTAAACAACATAAG	4079
4020	DB	CAGTTTGGAAATATTTACTTTCACAAGTATTGACACTGTTGTGGTATTAAACAACATAAG	4079
4080	QY	TTGCTCAAGGGCAATCATTTATTTCAAGTGGCTTAAAGTTTACTTCTGACAGTTTGTGTATA	4139
4080	DB	TTGCTCAAGGGCAATCATTTATTTCAAGTGGCTTAAAGTTTACTTCTGACAGTTTGTGTATA	4139
4140	QY	TTTATTTGGCTATTGGCAATTTGTTTTGTTTTTCTCTTTGGGTTTATTAATGTAAAGCA	4199
4140	DB	TTTATTTGGCTATTGGCAATTTGTTTTGTTTTTCTCTTTGGGTTTATTAATGTAAAGCA	4199
4200	QY	GGGATTTATTAACCTACAGTCCAGAAAGCCTGTGAATTTGAAATGAGGAAAAAATACATTT	4259
4200	DB	GGGATTTATTAACCTACAGTCCAGAAAGCCTGTGAATTTGAAATGAGGAAAAAATACATTT	4259
4260	QY	TTGTTTTTACCACTTCTAACTAAATTTAAACATTTTATTCCTATTCGGAATAGAGCCATAA	4319
4260	DB	TTATTTTACCACTTCTAACTAAATTTAAACATTTTATTCCTATTCGGAATAGAGCCATAA	4319
4320	QY	ACTCAAGTGGTAAATACAGTACCTGTGATTTTGTGCATTTACATAGAGAAATCAGACAT	4379
4320	DB	ACTCAAGTGGTAAATAGAGTACCTGTGATTTTGTGCATTTACATAGAGAAATCAGACAT	4379

QY 4380 TTATATATATATATACAGTTGTTCCAGATAGCTTGTAAAGTGAATATTTATATCTCAAACT 4439
DB 4380 TTATATATATATATACAGTTGTTCCAGTACGTTGTAAAGTGAATATTTATATCTCAAACT 4439
QY 4440 ACTTTGAAATATAGACCTCTGCTGGATCTGTTTAAACATATTAATAAAACATGTTTAA 4499
DB 4440 ACTTTGAAATATAGACCTCTGCTGGATCTGTTTAAACATATTAATAAAACATGTTTAA 4499
QY 4500 AATTTTGATATTTTGAT 4559
DB 4500 AATTTTGATATTTTGAT 4559
QY 4560 TATATATTTGAAACATCTTTCTGAGAAGATTTCCAGATTTTCCCAATGAGTTCTTG 4619
DB 4560 TATATATTTGAAACATCTTTCTGAGAAGATTTCCAGATTTTCCCAATGAGTTCTTG 4619
QY 4620 GCATGCACACACACAGATGAAAGTATGATTTAGAGCTAAACATTTGATGATGCTGAG 4679
DB 4620 GCATGCACACACACAGATGAAAGTATGATTTAGAGCTAAACATTTGATGATGCTGAG 4679
QY 4680 ATGCAAGACTGAAATAGAAAGTTCTCCCAAGATACACAGTTGTTTAAAGCTAGGGT 4739
DB 4680 ATGCAAGACTGAAATAGAAAGTTCTCCCAAGATACACAGTTGTTTAAAGCTAGGGT 4739
QY 4740 GAGGGGGGAAATCTGCGCTTCTATAGGAATGCTCTCCCTGGAGCTGTTGAGGCTGCT 4799
DB 4740 GAGGGGGGAAATCTGCGCTTCTATAGGAATGCTCTCCCTGGAGCTGTTGAGGCTGCT 4799
QY 4800 CCTTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4859
DB 4800 CCTTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4859
QY 4860 TGGATCTCCAGTTCTAGCATAGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4919
DB 4860 TGGATCTCCAGTTCTAGCATAGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 4919
QY 4920 GTGAATGGAATATTAAGTAAATATATATATATATATATATATATATATATATATATAT 4979
DB 4920 GTGAATGGAATATTAAGTAAATATATATATATATATATATATATATATATATATATAT 4979
QY 4980 GTCTAAGTGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5035
DB 4980 GTCTAAGTGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5039
QY 5036 AGATATAGGAATATTAAGTAAATATATATATATATATATATATATATATATATATATAT 5095
DB 5040 AGATATAGGAATATTAAGTAAATATATATATATATATATATATATATATATATATATAT 5099
QY 5096 AACTCCAAACAGACTTCTGGAAGTTATTTTCTAAGATCTTCTGCGAGCTGGAAGGCA 5155
DB 5100 AACTCCAAACAGACTTCTGGAAGTTATTTTCTAAGATCTTCTGCGAGCTGGAAGGCA 5159
QY 5156 ACCCCCTGTGCAGAGCCACCCAGCTCAGCTGCGGCACTCTGCTTCCCCCATGAAG 5215
DB 5160 ACCCCCTGTGCAGAGCCACCCAGCTCAGCTGCGGCACTCTGCTTCCCCCATGAAG 5219
QY 5216 GGCTGGCTCCCGATATATATAAACCCTCTCGAGCTCGGGCATGAGCCAGCAAGG 5271
DB 5220 GGCTGGCTCCCGATATATATAAACCCTCTCGAGCTCGGGCATGAGCCAGCAAGG 5275

RESULT 14
US-10-244-633-2
; Sequence 2, Application US/10244633
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis,
; TITLE OF INVENTION: Prognosis And Treatment Of Glaucoma And Related
; TITLE OF INVENTION: Disorders

FILE REFERENCE: 07425.0057.US01
; CURRENT APPLICATION NUMBER: US/10/244, 633
; CURRENT FILING DATE: 2002-09-17
; PRIOR APPLICATION NUMBER: US/09/306, 828
; PRIOR FILING DATE: 1999-05-07
; PRIOR APPLICATION NUMBER: US 09/227, 881
; PRIOR FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 2
; LENGTH: 5304
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-244-633-2

Query Match 99.1%; Score 5224.4; DB 43; Length 5304;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 5264; Conservative 0; Mismatches 6; Indels 6; Gaps 3;

QY 1 ATCTTTGTTTACGTTTACCTCAGGGCTATTATGAAATGAAATGAGATAACCAATGTGAAG 60
DB 1 ATCTTTGTTTACGTTTACCTCAGGGCTATTATGAAATGAAATGAGATAACCAATGTGAAG 60
QY 61 TCCTATAAACTGTATAGCTCCATTCCGATGTATCTTTGGCAGGATGATAAAGATCA 120
DB 61 TCCTATAAACTGTATAGCTCCATTCCGATGTATCTTTGGCAGGATGATAAAGATCA 120
QY 121 GGAAGAAGAGTATCCAGTTAGCCAAAGTGTCTCTTGGCAGGATGATAAAGATCA 180
DB 121 GGAAGAAGAGTATCCAGTTAGCCAAAGTGTCTCTTGGCAGGATGATAAAGATCA 180
QY 181 CAGATGTTGCTCTCCACAGAGCTATTCTTCAGGAAACATCACATCCAAATGTGTAATC 240
DB 181 CAGATGTTGCTCTCCACAGAGCTATTCTTCAGGAAACATCACATCCAAATGTGTAATC 240
QY 241 CATCAAAACAGGAGCTAAAGAAACAGGAATGAGATGGGCACTTCCCAAGGAAATGCCAG 300
DB 241 CATCAAAACAGGAGCTAAAGAAACAGGAATGAGATGGGCACTTCCCAAGGAAATGCCAG 300
QY 301 GAGAGCAAAATATGATGAAATTAACCTTTCCCTTTGTTTAAATTTTTCAGGAAATG 360
DB 301 GAGAGCAAAATATGATGAAATTAACCTTTCCCTTTGTTTAAATTTTTCAGGAAATG 360
QY 361 ATGAGGACCAAAATCAATGAATTAAGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
DB 361 ATGAGGACCAAAATCAATGAATTAAGAAACAGCTCAGAAAAAGATGTTTCCAAATGG 420
QY 421 TAATTAAGTATTTGTTTCTTGGGAAGAGACCTCCATGTGAGCTTGATGGGAAATGGAA 480
DB 421 TAATTAAGTATTTGTTTCTTGGGAAGAGACCTCCATGTGAGCTTGATGGGAAATGGAA 480
QY 481 AAACGTCAAAACATGATCTGATCAGATCCCAAGTGGATTTATTTTAAACCCAGAT 540
DB 481 AAACGTCAAAACATGATCTGATCAGATCCCAAGTGGATTTATTTTAAACCCAGAT 540
QY 541 GGCATCCTCTGGGAGGCAAGTTTCCAGGAAGTCTATGTTAGCAAAAGGACATTAACATAC 600
DB 541 GGCATCCTCTGGGAGGCAAGTTTCCAGGAAGTCTATGTTAGCAAAAGGACATTAACATAC 600
QY 601 AGCAAAATCAAAATTTCCGCAAAATGCAAGGAAAAATGGGAGCTGGGAAAGCTTTTCAAC 660
DB 601 AGCAAAATCAAAATTTCCGCAAAATGCAAGGAAAAATGGGAGCTGGGAAAGCTTTTCAAC 660
QY 661 AGTGATAGGCAAGTTGATCCATGTTGCAACACCTCCCGCTCTATACCAGGGAACACAAA 720
DB 661 AGTGATAGGCAAGTTGATCCATGTTGCAACACCTCCCGCTCTATACCAGGGAACACAAA 720
QY 721 ATTGATGGCTAAAGCTGAGCTTTCAAGGGAATATGAAAACTGAGAGCAAAACAAA 780
DB 721 ATTGATGGCTAAAGCTGAGCTTTCAAGGGAATATGAAAACTGAGAGCAAAACAAA 780
QY 781 GACATGTTTAAAGGCAACAGAACTTGTAGCCTTCAAGAGCAGTGGCCCTCAGCA 840

Db	781	GACATGGTTAAAGGCAACAGACATTTGTAGCCTTCAAAGCAGCAGTGCCTCCAGCA	840
Qy	841	GGGACCTTGAGCATTTCCTTTAGGAGCCAGTTTCTTAAGNACTTTAAGAACTC	900
Db	841	GGGACCTTGAGCATTTCCTTTAGGAGCCAGTTTCTTAAGNACTTTAAGAACTC	900
Qy	901	TTGAAAGATCATGAATTTTAAACATTTTAAAGTATAAACAATAATGCGATGATAATCAG	960
Db	901	TTGAAAGATCATGAATTTTAAACATTTTAAAGTATAAACAATAATGCGATGATAATCAG	960
Qy	961	TTTAGACATGGTCCCAATTTTATAAGTCAGGCATACAGGATAACGTGTCCAGCTCC	1020
Db	961	TTTAGACATGGTCCCAATTTTATAAGTCAGGCATACAGGATAACGTGTCCAGCTCC	1020
Qy	1021	GGATAGGTGAGAAATCATTAGAAATCACTGTGTCCCATCTCTAACTTTTTCAGAAATGATC	1080
Db	1021	GGATAGGTGAGAAATCATTAGAAATCACTGTGTCCCATCTCTAACTTTTTCAGAAATGATC	1080
Qy	1081	TGTCATAGCCTCACACAGGCCGATGTGTGACCTTACACACATCTACACCCAA	1140
Db	1081	TGTCATAGCCTCACACAGGCCGATGTGTGACCTTACACACATCTACACCCAA	1140
Qy	1141	GTGCTCAACCATTTGTTAAGTGTATCTCAGTAGGTCCCATTAACAATGCCACTCC	1200
Db	1141	GTGCTCAACCATTTGTTAAGTGTATCTCAGTAGGTCCCATTAACAATGCCACTCC	1200
Qy	1201	TGTGAGGCCATCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCATCAGATGT	1260
Db	1201	TGTGAGGCCATCCGCTCCACAGGAAGTCTCCCACTCTAGACTTCTGCATCAGATGT	1260
Qy	1261	TACAGCCAGAAAGTCCGTGAGGGTGAGGGTCTGTCTTACACCTACCTGTATGCTCTAC	1320
Db	1261	TACAGCCAGAAAGTCCGTGAGGGTGAGGGTCTGTCTTACACCTACCTGTATGCTCTAC	1320
Qy	1321	ACCTGAGTCACTGCAACCTCTGCTCCAGGTTTCAAGCAATTCCTCTGCTCAGCCTCC	1380
Db	1321	ACCTGAGTCACTGCAACCTCTGCTCCAGGTTTCAAGCAATTCCTCTGCTCAGCCTCC	1380
Qy	1381	CGCTGAGCTGGGACTACAGGCGCACGCGCGCTAATTTTGTATTTGTAGTAGAGTGGG	1440
Db	1381	CGCTGAGCTGGGACTACAGGCGCACGCGCGCTAATTTTGTATTTGTAGTAGAGTGGG	1440
Qy	1441	GTTCACCATATTAGCCCGCTGGTCTTTGAACCTCTGACCTCAGGTATCACCCACCTC	1500
Db	1441	GTTCACCATATTAGCCCGCTGGTCTTTGAACCTCTGACCTCAGGTATCACCCACCTC	1500
Qy	1501	AGCTCTTAAGTGTGGGATTAAGGCATGAGTCAACGCGCCCGGCAAGGTCAGTGT	1560
Db	1501	AGCTCTTAAGTGTGGGATTAAGGCATGAGTCAACGCGCCCGGCAAGGTCAGTGT	1560
Qy	1561	TTAATAAGGAATACTTGAATGGTTTACTAAACCAACAGGGAACACAGCAAAAGCTCTGA	1620
Db	1561	TTAATAAGGAATACTTGAATGGTTTACTAAACCAACAGGGAACACAGCAAAAGCTCTGA	1620
Qy	1621	TAAITTCAGGAACTTTTGGGATGGGGAATGGTGCATGAGCTGCTGCTAGTCCAGAC	1680
Db	1621	TAAITTCAGGAACTTTTGGGATGGGGAATGGTGCATGAGCTGCTGCTAGTCCAGAC	1680
Qy	1681	CAGTGTCTCATCAGCTTTCTTCCCTCATCTCTGATTTTTCAGGCTAAGTTACATTTATT	1740
Db	1681	CAGTGTCTCATCAGCTTTCTTCCCTCATCTCTGATTTTTCAGGCTAAGTTACATTTATT	1740
Qy	1741	CACCATGCTTTTGTGGTAAAGCTCCACATCGTTACTGAATTAAGAGTATACATAAATAG	1800
Db	1741	CACCATGCTTTTGTGGTAAAGCTCCACATCGTTACTGAATTAAGAGTATACATAAATAG	1800
Qy	1801	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGGAGGGCATACCCAGAGACTCCT	1860
Db	1801	TTCCATTTGGGGCCATCTGTGTGTGTATAGGGAGGAGGGCATACCCAGAGACTCCT	1860
Qy	1861	TGAAGCCCCGGCAGAGGTTCTCTCAGTGGGGAGGCCCTGCAAGACACCGGGGTCC	1920
Db	1861	TGAAGCCCCGGCAGAGGTTCTCTCAGTGGGGAGGCCCTGCAAGACACCGGGGTCC	1920
Qy	1921	TGGGTGCTCTGAGCAACCTGCGACCGCTGCGCACTGGTGTGTTTGTATCACTCTCTAGG	1980
Db	1921	TGGGTGCTCTGAGCAACCTGCGACCGCTGCGCACTGGTGTGTTTGTATCACTCTCTAGG	1980
Qy	1981	GACCTGTTGCTTTCTATTTCTGTGTGACTCGTTCATTCATCCAGGCATTCATTGACAAT	2040
Db	1981	GACCTGTTGCTTTCTATTTCTGTGTGACTCGTTCATTCATCCAGGCATTCATTGACAAT	2040
Qy	2041	TATTGAGTACTTATATCTGCCAGACACAGAGACAAATATGCTGAGCAAGAGTCACTGC	2100
Db	2041	TATTGAGTACTTATATCTGCCAGACACAGAGACAAATATGCTGAGCAAGAGTCACTGC	2100
Qy	2101	CCTACCTTCGTGGAGGTGACAGTTTCTCATCGAAGACGTGAGAGAAATTAATAGCCA	2160
Db	2101	CCTACCTTCGTGGAGGTGACAGTTTCTCATCGAAGACGTGAGAGAAATTAATAGCCA	2160
Qy	2161	GCCAACTTAAACCCAGTGTCTGAAAGAAAGGAAATAAACCATCTCTTGAAGAAATTTGTGGC	2220
Db	2161	GCCAACTTAAACCCAGTGTCTGAAAGAAAGGAAATAAACCATCTCTTGAAGAAATTTGTGGC	2220
Qy	2221	AGCATCCCTTAAAGAGGCCACCTCCTAGCGCCCTCTGCTGCTCCATCTGTCGCCGAGG	2280
Db	2221	AGCATCCCTTAAAGAGGCCACCTCCTAGCGCCCTCTGCTGCTCCATCTGTCGCCGAGG	2280
Qy	2281	CCCCAAGCCGAGTCTTCAAGCCTCTCTCCATCAGTCACAGCGCTGAGCTGGCT	2340
Db	2281	CCCCAAGCCGAGTCTTCAAGCCTCTCTCCATCAGTCACAGCGCTGAGCTGGCT	2340
Qy	2341	GCCTCGCTTCCGCTGAAATCGTCTGCTGTCATCTGAGCTGGAGACTCTCTTGGCTCCAGGCT	2400
Db	2341	GCCTCGCTTCCGCTGAAATCGTCTGCTGTCATCTGAGCTGGAGACTCTCTTGGCTCCAGGCT	2400
Qy	2401	CCAGAAAGGAAATGAGAGGGAATCTAGTCTAACCGAGAAATCTGAGAGGGAACAGTGTTC	2460
Db	2401	CCAGAAAGGAAATGAGAGGGAATCTAGTCTAACCGAGAAATCTGAGAGGGAACAGTGTTC	2460
Qy	2461	CTCAGAGGGAAGAGGGCTCCACGTCAGAGAAATCCAGAGGTGGGAGTCCAGGGAG	2520
Db	2461	CTCAGAGGGAAGAGGGCTCCACGTCAGAGAAATCCAGAGGTGGGAGTCCAGGGAG	2520
Qy	2521	TGGGAGCGCTGGGCTGAGCGGTGCTGAAAGGAGGGAAGGTGAAAGGGCAAGGCTGAA	2580
Db	2521	TGGGAGCGCTGGGCTGAGCGGTGCTGAAAGGAGGGAAGGTGAAAGGGCAAGGCTGAA	2580
Qy	2581	GCTGCCAGATGTTTCAAGTGTGTTTACGCGGCTGGGAGTTCCTGTTCTCTGTGAGC	2640
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Qy	2641	CTTTTATCTTTTCTCTGCTTGGAGAGAGTCTATTTCTATGAAGGATGAGTTC	2700
Db	2641	CTTTTATCTTTTCTCTGCTTGGAGAGAGTCTATTTCTATGAAGGATGAGTTC	2700
Qy	2701	ATAAGTCAGCTGTTAAATTTCCAGGGTGTGCAATGGGTTTCTTCCATCAAGAGGCTTTAT	2760
Db	2701	ATAAGTCAGCTGTTAAATTTCCAGGGTGTGCAATGGGTTTCTTCCATCAAGAGGCTTTAT	2760
Qy	2761	TAAATGGGAATATAGGAAGCGAGCTCATTTCTTAGGCGGTAAATTAACGGAAGTGCAC	2820
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Qy	2821	TGGAGTCTTTTCTTCTCATGTCTTCTGCGCAACTACTCAGCCCTGCTGAGCTTGGCTTA	2880
Db	2821	TGGAGTCTTTTCTTCTCATGTCTTCTGCGCAACTACTCAGCCCTGCTGAGCTTGGCTTA	2880
Qy	2881	TGCAAGACGCTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTCTTCTTCTGCAAT	2940
Db	2881	TGCAAGACGCTCGAAACCTTGGAAATCAGGAGACTCGGTTTCTTCTTCTTCTGCAAT	2940
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Db	2941	GGTTGCTGTGGACCGTGGGCAAGTGTCTCTCTTCCCTGGGCCATAGTCTTCTCTGCT	3000

QY	3001	ATAAAGACCCCTTGCAGCTCTCGTGTCTGTGAAACATCTCCCTGTGATCTCTCTGTGAGGG	3060
Db	3001	TTGCTCAAAGGCAATCAATATATTTCAAGTGGCTTAAAGTTACTCTCTGACAGTTTGTGTATA	4133
QY	3061	GGATGTTGAGAGGGGAAGGAGCGCAGCTGTGAGCGAGCTGTAGCCACAGGGAGGTGTGAGGG	3120
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QY	3121	GGACAGGAAGGAGCGCAGAAAGCTGGGTGTCTCCATCAGTCCCTCACTGTATCAGCTCAGACTC	3180
Db	3121	GGATTTATTAACCTTACAGTCCAGAAAGCCCTGTGAATTTGAATGAGGAAAAATTAAGTTT	4259
QY	3181	CAGGACCGGAGAGCCCAATATGCTTTCAGGAAAGCTCAATGAACCCAAACGACCAATTTTCT	3240
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QY	3301	GSTAGCTTTTGGCTGGCATTTCAAAACATGGGCGAGCAAGTGGAAAAATGCCAGAGATTG	3360
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QY	3361	TTAAACTTTTCACTTGAACGACCCCGAGCTCAGCAGTCTCAGCAGTCTCAGCAGCACTG	3420
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QY	3421	AGTGACTCTGAGCGGAGGAGGAGNAGAAAAAGAGAGGGATAGTGTATGAGCAAGAAAG	3480
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QY	3721	GGAGTTAGCAGCAAGAGGCAATCCGTTTCTTTTAAACGAGGAAGAAACATTCCTAAGAG	3780
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DB	2461	CTCAGAGGAAAGGGGCTTCCAGCTCCAGGAAATCCAGAGGTGGGGA	CTCAGGGAG	2520
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OM nucleic - nucleic search, using sw model

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Minimum DB seq length: 0

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Post-processing: Minimum Match 0%

Maximum Match 100%
Listing first 45 summaries

Database : Pending Patents NA New: *

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Database : Pending_Patents_NA_New:*
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2: /cgn2_6/ptodata/1/pna/US06 NEW COMB.seq:*
3: /cgn2_6/ptodata/1/pna/US07 NEW COMB.seq:*
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12: /cgn2_6/ptodata/1/pna/US60 NEW COMB.seq:*
17Jan05:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	184.8	3.5	1000	11	US-11-266-748A-198985	Sequence 198985,
C 2	184.4	3.5	1000	11	US-11-266-748A-224089	Sequence 224089,
C 3	184.4	3.5	1770	11	US-11-266-748A-26905	Sequence 26905, A
4	184.2	3.5	1000	11	US-11-266-748A-195297	Sequence 195297,
5	183.6	3.5	17691	6	US-10-868-184C-6721	Sequence 6721, Ap
6	183.6	3.5	17691	7	US-10-868-184C-6721	Sequence 6721, Ap
7	182.4	3.5	515	11	US-11-266-748A-212844	Sequence 212844,
C 8	182.4	3.5	515	11	US-11-266-748A-236144	Sequence 236144,
C 9	180.8	3.4	101500	11	US-11-266-748A-61740	Sequence 61740, A
10	180.6	3.4	39729	11	US-11-214-063A-1285	Sequence 1285, Ap
11	179.4	3.4	10948	11	US-11-214-063A-1715	Sequence 1715, Ap
C 12	179.2	3.4	6708	6	US-10-868-184C-6360	Sequence 6360, Ap
C 13	179.2	3.4	6708	6	US-10-868-184C-11367	Sequence 11367, A
C 14	179.2	3.4	6708	7	US-10-868-184C-6360	Sequence 6360, Ap
C 15	179.2	3.4	6708	7	US-10-868-184C-11367	Sequence 11367, A
C 16	179.2	3.4	189450	1	PCT-US05-10912-6	Sequence 6, Appli
C 17	179.2	3.4	495475	11	US-11-266-748A-28223	Sequence 28223, A
C 18	178.6	3.4	124048	11	US-11-266-748A-53707	Sequence 53707, A
C 19	177.8	3.4	153379	11	US-11-266-748A-24170	Sequence 24170, A
C 20	177.8	3.4	155379	11	US-11-266-748A-59410	Sequence 59410, A
C 21	177.8	3.4	227968	11	US-11-266-748A-60135	Sequence 60135, A
C 22	177.8	3.4	227968	11	US-11-266-748A-60135	Sequence 60135, A

C 22	177.6	3.4	128978	11	US-11-2666-748A-61436	Sequence 61436, A
C 23	176.8	3.4	53332	6	US-10-786-065-3	Sequence 3, Appli
C 24	176.8	3.4	53332	7	US-10-786-065-3	Sequence 3, Appli
C 25	176.6	3.4	1173	6	US-10-461-673-4221	Sequence 4221, Ap
C 26	176.6	3.4	1173	8	US-10-461-673-4221	Sequence 4221, Ap
C 27	176	3.3	1400	6	US-10-868-184C-7974	Sequence 7974, Ap
C 28	176	3.3	1400	7	US-10-868-184C-7974	Sequence 7974, Ap
C 29	176	3.3	2213	6	US-10-868-184C-7973	Sequence 7973, Ap
C 30	176	3.3	2213	7	US-10-868-184C-7973	Sequence 7973, Ap
C 31	176	3.3	164429	11	US-11-2666-748A-22662	Sequence 22662, A
C 32	175.6	3.3	1000	11	US-11-2666-748A-196932	Sequence 196932, A
C 33	175.6	3.3	3143	11	US-11-2666-748A-28298	Sequence 28298, A
C 34	175.4	3.3	6110	11	US-11-2666-748A-24483	Sequence 24483, A
C 35	175.4	3.3	100086	11	US-11-2666-748A-5715	Sequence 5715, A
C 36	175.2	3.3	65854	6	US-10-868-184C-12628	Sequence 12628, A
C 37	175.2	3.3	65854	6	US-10-868-184C-13035	Sequence 13035, A
C 38	175.2	3.3	65854	7	US-10-868-184C-12628	Sequence 12628, A
C 39	175.2	3.3	65854	7	US-10-868-184C-13035	Sequence 13035, A
C 40	175.2	3.3	832900	11	US-11-214-063A-13313	Sequence 13313, Ap
C 41	175	3.3	1000	11	US-11-2666-748A-197285	Sequence 197285, A
C 42	175	3.3	2297	6	US-10-276-817B-3434	Sequence 3434, Ap
C 43	175	3.3	2297	7	US-10-276-817B-3434	Sequence 3434, Ap
C 44	175	3.3	6615	11	US-11-2666-748A-30008	Sequence 30008, A
C 45	175	3.3	6639	11	US-11-2666-748A-30006	Sequence 30006, A

ALIGNMENTS

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RESULT 1
US-11-266-748A-198985
; Sequence 198985, Application US/11266748A
; GENERAL INFORMATION:
; APPLICANT: Harkin, Paul
; APPLICANT: Johnston, Patrick
; APPLICANT: Mulligan, Karl
; TITLE OF INVENTION: Transcriptome Microarray Technology and
; TITLE OF INVENTION: Methods of Using the Same
; FILE REFERENCE: 55815-0102 (319189)
; CURRENT APPLICATION NUMBER: US/11/266,748A
; CURRENT FILING DATE: 2005-11-03
; PRIOR APPLICATION NUMBER: EP 04105479.2
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105482.6
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105483.4
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105507.0
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105507.0
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105485.9
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105484.2
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: US 60/662,276
; PRIOR FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/700,293
; PRIOR FILING DATE: 2005-07-18
; NUMBER OF SEQ ID NOS: 483996
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 198985
; LENGTH: 1000
; TYPE: DNA
; ORGANISM: Homo Sapiens
US-11-266-748A-198985

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Query Match	3.5%;	Score 184.8;	DB 11;	Length 1000;
Best Local Similarity	78.2%;	Pred. No. 5.3e-36;		
Matches 222;	Conservative 0;	Mismatches 62;	Indels 0;	Gaps 0;
1281	GGGTGAGGGTCTGTGTTCTTTACACCTGATGTGTCTACACCTGAGTCTCACTGCAACCT			1340
578	SAGTCTCGCTCTGTTACCTACCATCTGGAGTGCAGTGGCGCTATCTCAGCTCACTGCAACCT			637

QY 1341 CTCCTCCAGGTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGACTACAGG 1400
DB 638 CCGCTCCCGGTTTCAAGCAATCTCTGTCTCAGCTCCCGGTAGCTGGACTACAGG 697
QY 1401 CGCAGCCCGGCTAAATTTTGTATGTTAGTAGAGTGGGTTTCCACATATTAGCCCGG 1460
DB 698 CTCATGCCAATTTAAATTTTGTATGTTAGTAGAGTGGGTTTCCCGGTAGCTGG 757
QY 1461 CTGTCTTGAACCTCTGACCTCAGGTGATCCACCCAGCTCAGCCCTCTAAAGTCTGGGA 1520
DB 758 CTGTCTTGAACCTCTGACCTCAGGTGATCCACCCAGCTCAGCCCTCTAAAGTCTGGGA 817
QY 1521 TTACAGCATGAGTCACGGCCCGGCAAGGGTCAGTGTAA 1564
DB 818 TTACAGCATGAGTCACGGCCCGGCAAGGGTCAGTGTAA 861

RESULT 2
US-11-266-748A-224089/c
; Sequence 224089, Application US/11266748A
; GENERAL INFORMATION:
; APPLICANT: Harkin, Paul
; APPLICANT: Mulligan, Karl
; APPLICANT: Johnston, Patrick
; TITLE OF INVENTION: Transcriptome Microarray Technology and
; FILE REFERENCE: 55815-0102 (319189)
; CURRENT FILING DATE: 2005-11-03
; PRIOR APPLICATION NUMBER: EP 04105479.2
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105482.6
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105483.4
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105507.0
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105485.9
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105485.9
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105484.2
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: US 60/662,276
; PRIOR FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/700,293
; PRIOR FILING DATE: 2005-07-18
; NUMBER OF SEQ ID NOS: 483996
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 224089
; LENGTH: 1000
; TYPE: DNA
; ORGANISM: Homo Sapiens
US-11-266-748A-224089

Query Match 3.5%; Score 184.4; DB 11; Length 1000;
Best Local Similarity 81.0%; Pred. No. 6.7e-36;
Matches 230; Conservative 0; Mismatches 46; Indels 8; Gaps 1;
QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 1380
DB 938 ATCTCAGCTCACTGCAACCTCTGCTCCAGGTTCAAGGATTTCCCTGCTCAGCTCC 879
QY 1381 CGCGTAGCTGGGACTACAGGCG- - - - -CAGCGCCGGCTAAATTTTGTATTGTAGTA 1432
DB 878 CGAGCAGCTGGGACTACAGGCGCCGACACACCGCGGCTAAATTTTGTATTGTAGTA 819
QY 1433 GAGATGGGTTTCCACCATATTAGCCCGGCTGTCTTGAACTCCTGACCTCAGGTGATCCA 1492
DB 818 GAGATGGGTTTCCACCATATTAGCCCGGCTGTCTTGAACTCCTGACCTCAGGTGATCTG 759
QY 1493 CCCACCTCAGCTCTTAAAGTCTGGGATTTACAGGATGAGTCACCGCGCCGCGCAAGG 1552
DB 758 CCCGCTCGGCTCTCCAAAGTGGGATTTACAGGATGAGCCACCGCATCCGGCCAGAT 699

QY 1553 GTCAGTGTTTAATAAGGATTAATGATGTTTACTTAACCAA 1596
DB 698 TTCAGGTGCTTTTAAAGAAAGTAAGTGAATTTTATTTTACTTAA 655
RESULT 3
US-11-266-748A-26905/c
; Sequence 26905, Application US/11266748A
; GENERAL INFORMATION:
; APPLICANT: Harkin, Paul
; APPLICANT: Johnston, Patrick
; APPLICANT: Mulligan, Karl
; TITLE OF INVENTION: Transcriptome Microarray Technology and
; FILE REFERENCE: 55815-0102 (319189)
; CURRENT FILING DATE: 2005-11-03
; PRIOR APPLICATION NUMBER: EP 04105479.2
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105482.6
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105483.4
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105507.0
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105485.9
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: EP 04105484.2
; PRIOR FILING DATE: 2004-11-03
; PRIOR APPLICATION NUMBER: US 60/662,276
; PRIOR FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/700,293
; PRIOR FILING DATE: 2005-07-18
; NUMBER OF SEQ ID NOS: 483996
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 26905
; LENGTH: 1770
; TYPE: DNA
; ORGANISM: Homo Sapiens
US-11-266-748A-26905

Query Match 3.5%; Score 184.4; DB 11; Length 1770;
Best Local Similarity 81.0%; Pred. No. 8.4e-36;
Matches 230; Conservative 0; Mismatches 46; Indels 8; Gaps 1;
QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 1380
DB 1708 ATCTCAGCTCACTGCAACCTCTGCTCCAGGTTCAAGGATTTCCCTGCTCAGCTCC 1649
QY 1381 CGCGTAGCTGGGACTACAGGCG- - - - -CAGCGCCGGCTAAATTTTGTATTGTAGTA 1432
DB 1648 CGAGCAGCTGGGACTACAGGCGCCGACACACCGCGGCTAAATTTTGTATTGTAGTA 1589
QY 1433 GAGATGGGTTTCCACCATATTAGCCCGGCTGTCTTGAACTCCTGACCTCAGGTGATCCA 1492
DB 1588 GAGATGGGTTTCCACCATATTAGCCCGGCTGTCTTGAACTCCTGACCTCAGGTGATCTG 1529
QY 1493 CCCACCTCAGCTCTTAAAGTCTGGGATTTACAGGATGAGTCACCGCGCCGCGCAAGG 1552
DB 1528 CCCGCTCGGCTCTCCAAAGTGGGATTTACAGGATGAGCCACCGCATCCGGCCAGAT 1469
QY 1553 GTCAGTGTTTAATAAGGATTAATGATGTTTACTTAACCAA 1596
DB 1468 TTCAGGTGCTTTTAAAGAAAGTAAGTGAATTTTATTTTACTTAA 1425

RESULT 4
US-11-266-748A-195297
; Sequence 195297, Application US/11266748A
; GENERAL INFORMATION:
; APPLICANT: Harkin, Paul
; APPLICANT: Johnston, Patrick
; APPLICANT: Mulligan, Karl

[illegible]

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RES001 5
US/10-868-184C-6721
; Sequence 6721, Application US/10868184C
; GENERAL INFORMATION:
; APPLICANT: Rosen, et. al
; TITLE OF INVENTION: Human Secreted Proteins
; FILE REFERENCE: PS905
; CURRENT APPLICATION NUMBER: US/10/868,184C
; CURRENT FILING DATE: 2004-06-16
; PRIOR APPLICATION NUMBER: 60/278,650
; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: 09/833,245
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US01/11988
; PRIOR FILING DATE: 2001-04-12

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PRIOR APPLICATION NUMBER: PCT/US00/06042
PRIOR FILING DATE: 2000-03-09
BEST LOCAL SIMILARITY: PCT/US00/06014
PRIOR FILING DATE: 2000-03-09
REMAINING PRIOR APPLICATION DATA REMOVED - SEE FILE WRAPPER OR PALM.
NUMBER OF SEQ ID NOS: 13046
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6721
LENGTH: 17691
TYPE: DNA
ORGANISM: Homo sapiens
US-10-868-184C-6721

Query Match 3.5%; Score 183.6; DB 7; Length 17691;
Best Local Similarity 78.2%; Pred. No. 3.4e-35;
Matches 233; Conservative 0; Mismatches 64; Indels 1; Gaps 1;

QY 1283 GTGAGGCTGTGCTTACACCTACCTGCTATGCTCTACACCTGAGTCACTGCAACCTCT 1342
DB 14938 GTCTCGCTCTGTCCGCCAGGAGGAGTGCAATGCGCTCAGCTCAGCTGCAACCTCT 14997

QY 1343 GCCTCCAGGTTCAAGCAATCTCTGTCTCAGCCTCCCGCTAGCTGGGACTACAGCG 1402
DB 14998 GCCTCCAGGTTCAAGCAATCTCTGTCTCAGCCTCCCGCTAGCTGGGACTACAGCG 15057

QY 1403 CAGCCCGCGCTAA-TTTTTGATTTAGTAGATGGGTTTACCATATTAGCCCGC 1461
DB 15058 TGCGCCAGCTAAATTTTGTATTTTAGTAGACAGAGGTTTACCATATTAGCCCGC 15117

QY 1462 TGGCTTGAACCTCGACCTCAGGTGATCCACCACTCAGCTCCTAAAGTGTGGAT 1521
DB 15118 TGGCTTGAACCTCGACCTCAGGTGATCCACCACTCAGCTCCTAAAGTGTGGAT 15177

QY 1522 TACAGGATGATGATCAGCGCGCCGCGCAAGGTCAGTGTGTTAATAGGAATACTCA 1579
DB 15178 TATAGGATGAGCAGCGTGTCCGCCAGAGGCTTCTTTAAGTGTGACTTCCAA 15235

RESULT 7
US-11-266-748A-212844
Sequence 212844, Application US/11266748A
GENERAL INFORMATION:
APPLICANT: Harkin, Paul
APPLICANT: Mulligan, Karl
TITLE OF INVENTION: Transcriptome Microarray Technology and
TITLE OF INVENTION: Methods of Using the Same
FILE REFERENCE: 55815-0102 (319189)
CURRENT APPLICATION NUMBER: US/11/266,748A
PRIOR FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: EP 04105479.2
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105482.6
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105483.4
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105507.0
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105485.9
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105484.2
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: US 60/662,276
PRIOR FILING DATE: 2005-03-14
PRIOR APPLICATION NUMBER: US 60/700,293
PRIOR FILING DATE: 2005-07-18
NUMBER OF SEQ ID NOS: 483996
SOFTWARE: PatentIn version 3.3
SEQ ID NO 212844
LENGTH: 515
TYPE: DNA
ORGANISM: Homo sapiens
US-11-266-748A-212844

Query Match 3.5%; Score 182.4; DB 11; Length 515;
Best Local Similarity 86.4%; Pred. No. 1.6e-35;
Matches 216; Conservative 0; Mismatches 26; Indels 8; Gaps 1;

QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 1380
DB 46 ATCTCAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 105

QY 1381 CGCTAGCTGGGACTACAGCG-----CAGCCCGGCTAATTTTGTATTGTAGTA 1432
DB 106 CAAGTAGCTGGGATTACAGCGCACACCAACCGCCGGCTAATTTTGTATTGTAGTA 165

QY 1433 GAGATGGGTTTACCATTATTAGCCCGGCTGCTTGAACCTCTGACCTCAGCTGATCCA 1492
DB 166 GAGATGGGTTTACCATTATTAGCCCGGCTGCTTGAACCTCTGACCTCAGCTGATCCA 225

QY 1493 CCACCTCAGCTCTTAAAGTGTGGGATTACAGGATGAGTCAACCGCGCCCGCAAGG 1552
DB 226 CCACCTCGGCTCCCAAGTGTGGGATTACAGGCTGGGCCACTGGCCCACTTAAT 285

QY 1553 GTCAGTGT 1562
DB 286 TTTGTATT 295

RESULT 8
US-11-266-748A-236144/c
Sequence 236144, Application US/11266748A
GENERAL INFORMATION:
APPLICANT: Harkin, Paul
APPLICANT: Johnston, Patrick
APPLICANT: Mulligan, Karl
TITLE OF INVENTION: Transcriptome Microarray Technology and
TITLE OF INVENTION: Methods of Using the Same
FILE REFERENCE: 55815-0102 (319189)
CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: EP 04105479.2
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105482.6
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105483.4
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105507.0
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105485.9
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105484.2
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: US 60/662,276
PRIOR FILING DATE: 2005-03-14
PRIOR APPLICATION NUMBER: US 60/700,293
PRIOR FILING DATE: 2005-07-18
NUMBER OF SEQ ID NOS: 483996
SOFTWARE: PatentIn version 3.3
SEQ ID NO 236144
LENGTH: 515
TYPE: DNA
ORGANISM: Homo sapiens
US-11-266-748A-236144

Query Match 3.5%; Score 182.4; DB 11; Length 515;
Best Local Similarity 86.4%; Pred. No. 1.6e-35;
Matches 216; Conservative 0; Mismatches 26; Indels 8; Gaps 1;

QY 1321 ACCTGAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 1380
DB 470 ATCTCAGCTCACTGCAACCTCTGCTCCAGGTTCAAGCAATTTCTCTGTCTCAGCTCC 411

QY 1381 CGCTAGCTGGGACTACAGCG-----CAGCCCGGCTAATTTTGTATTGTAGTA 1432
DB 410 CAAGTAGCTGGGATTACAGCGCACACCAACCGCCGGCTAATTTTGTATTGTAGTA 351

QY 1433 GAGATGGGTTTACCATATTAGCCGGCTGGTCTTGAATCTCTGACCTCAGTGTATCCA 1492
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 350 GAGATGGGTTTACCATATTAGCCAGGCTGGTCTGAACTCTGACCTCAGTGTATCCA 291
 QY 1493 CCACCTCAGCTCTTAAGTGTCTGGATACAGGCATGAGTACCGCCCGGCCAAGG 1552
 Db |||||
 290 CCACCTCAGCTCTCCCAAGTGTCTGGATACAGGCCTGGCCACTGCGCCAGCCTAAT 231
 QY 1553 GTCAGTGT 1562
 Db |||||
 230 TTTTGTATTT 221

RESULT 9
 US-11-266-748A-61740
 ; Sequence 61740, Application US/11266748A
 ; GENERAL INFORMATION:
 ; APPLICANT: Harkin, Paul
 ; APPLICANT: Johnston, Patrick
 ; APPLICANT: Mulligan, Karl
 ; TITLE OF INVENTION: Transcriptome Microarray Technology and
 ; FILE REFERENCE: Methods of Using the Same
 ; CURRENT APPLICATION NUMBER: US/11/266,748A
 ; CURRENT FILING DATE: 2005-11-03
 ; PRIOR APPLICATION NUMBER: EP 04105479.2
 ; PRIOR FILING DATE: 2004-11-03
 ; PRIOR APPLICATION NUMBER: EP 04105482.6
 ; PRIOR FILING DATE: 2004-11-03
 ; PRIOR APPLICATION NUMBER: EP 04105483.4
 ; PRIOR FILING DATE: 2004-11-03
 ; PRIOR APPLICATION NUMBER: EP 04105507.0
 ; PRIOR FILING DATE: 2004-11-03
 ; PRIOR APPLICATION NUMBER: EP 04105485.9
 ; PRIOR FILING DATE: 2004-11-03
 ; PRIOR APPLICATION NUMBER: EP 04105484.2
 ; PRIOR FILING DATE: 2004-11-03
 ; PRIOR APPLICATION NUMBER: US 60/662,276
 ; PRIOR FILING DATE: 2005-03-14
 ; PRIOR APPLICATION NUMBER: US 60/700,293
 ; PRIOR FILING DATE: 2005-07-18
 ; NUMBER OF SEQ ID NOS: 483996
 ; SOFTWARE: PatentIn version 3.3
 ; SEQ ID NO 61740
 ; LENGTH: 101500
 ; TYPE: DNA
 ; ORGANISM: Homo Sapiens
 ; US-11-266-748A-61740

Query Match 3.4%; Score 180.8; DB 11; Length 101500;
 Best Local Similarity 79.0%; Pred. No. 3.4e-34;
 Matches 215; Conservative 0; Mismatches 57; Indels 0; Gaps 0;
 QY 1281 GGGTCAGGCTCTGTCTTTACACCTACCTGTATGCTTACACCTGAGCTCACTGCAACCT 1340
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 76041 GAGTCTCGCTCTGTTGCCAGCTAGAGTGCAATGCGCATCTTGGCTCATTCGCACT 76100
 QY 1341 CTGCTCCAGGTTCAAGCAATTCCTGTCTTACGCTCCCGGTAGCTGGGACTACAGG 1400
 Db |||||
 76101 CGGCTCCAGGTTCAAGCGATTCCTCTGCTCAGTCACTAGTAGCTGGGATACAGG 76160
 QY 1401 CGCAGCCCGGCTAATTTTGTATTTAGTAGAGATGGGTTTACCATATTAGCCCGG 1460
 Db |||||
 76161 CATGACCCCGGCTAATTTTGTATTTAGTAGAGACAGAGTTTACCATATTGGCCAGG 76220
 QY 1461 CTGCTCTTGAATCTCTGACCTCAGGTGATCCACCCACCTCAGCTCTTAAAGTGTGGGA 1520
 Db |||||
 76221 CTGGTCTGAACTCTGACCTCAGGTGATCCACCCACCTCAGCTCTCCAAAGTGTGGGA 76280
 QY 1521 TTACAGGCATGAGTACCGGCCCGGCCAAGG 1552
 Db |||||
 76281 TTACAGGCGTGAGGCACTGTGCCCGGCCCAAG 76312

RESULT 10
 US-11-214-063A-1285
 ; Sequence 1285, Application US/11214063A
 ; GENERAL INFORMATION:
 ; APPLICANT: SUWA, MAKIKO
 ; APPLICANT: ASAI, KIYOSHI
 ; APPLICANT: AKIYAMA, YUTAKA
 ; APPLICANT: ABURATANI, HIROYUKI
 ; TITLE OF INVENTION: GUANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
 ; FILE REFERENCE: 084335/166
 ; CURRENT APPLICATION NUMBER: US/11/214,063A
 ; CURRENT FILING DATE: 2005-08-30
 ; PRIOR APPLICATION NUMBER: US/10/292,798
 ; PRIOR FILING DATE: 2002-11-13
 ; PRIOR APPLICATION NUMBER: 10/017,161
 ; PRIOR FILING DATE: 2001-12-18
 ; PRIOR APPLICATION NUMBER: JP 2001-246789
 ; PRIOR FILING DATE: 2001-06-18
 ; NUMBER OF SEQ ID NOS: 2070
 ; SOFTWARE: PatentIn Ver. 2.1
 ; SEQ ID NO 1285
 ; LENGTH: 39729
 ; TYPE: DNA
 ; ORGANISM: Homo sapiens
 ; FEATURE:
 ; LOCATION: source
 ; FEATURE:
 ; LOCATION: (1)..(39729)
 ; NAME/KEY: CDS
 ; LOCATION: (201)..(409)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (1705)..(1966)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (3861)..(4281)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (6128)..(6513)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (13450)..(13658)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (15064)..(15254)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (15941)..(16150)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (20045)..(20170)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (22124)..(22422)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (22512)..(22683)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (23437)..(23604)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (24073)..(24309)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (27352)..(27646)
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: (28263)..(28408)
 ; FEATURE:

Db 1481 GCTCGCTCAACCCAGGCTGGAGTGTAGTGGCGGAGTCTCAGCTCACTGCAACCTCTGCTC 1540
Qy 1348 CAGAGTTCAAGCAATTTCTCTGCTCTCAGCTCCCGGTAGCTGGGACTACAGCGCACGC 1407
Db 1541 CTGGGTTCAACAATTTCTCTGCTCAGCTCCCAAGTACAGTGGGATTAAGTGTGTAC 1600
Qy 1408 CCGGCTAAATTTTGTATTTAGTGTAGAGATGGGGTTTCAACATATTAGCCCGCTGGTCT 1467
Db 1601 CCGGCTAAATTTTGTATTTAGTGGAGATGGGGTTTCAACATATTAGCCCGCTAGTCT 1660
Qy 1468 TGAACCTCTGACCTCAGGTGATCCACCACTCAGCTCTCTAAAGTGTGGGATTAACAG 1527
Db 1661 TGAACCTCTGACCTCAGGTGATCCACCACTCAGCTCTCTAAAGTGTGGGATTAACAG 1720
Qy 1528 CATGAGTCAACCGCGCCGCCAAGGTCAGTGTGTTAA 1564
Db 1721 CATGAGCACTGCACCAAGACCCCACTCAGTGTGTTAA 1757

RESULT 12

US-10-868-184C-6360/c
; Sequence 6360, Application US/10868184C
; GENERAL INFORMATION:

; APPLICANT: Rosen, et. al
; TITLE OF INVENTION: Human Secreted Proteins
; FILE REFERENCE: PS805
; CURRENT APPLICATION NUMBER: US/10/868,184C
; PRIOR FILING DATE: 2004-06-16
; PRIOR APPLICATION NUMBER: 60/278,650
; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: 09/833,245
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US01/11988
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US00/06043
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06012
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06058
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06044
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06059
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06042
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06014
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13046
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 6360
; LENGTH: 6708
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-868-184C-6360

Query Match 3.4%; Score 179.2; DB 6; Length 6708;
Best Local Similarity 77.9%; Pred. No. 2.9e-34;
Matches 232; Conservative 0; Mismatches 58; Indels 8; Gaps 1;
Qy 1290 TCTGTGCTTACACCTACCTCTGATGCTCTACACCTGAGCTCACTGCAACCTCTGCTCCC 1349
Db 3909 TCTGTCTCTCAGGCTGGAGTACAGTGGGCACAACTCTCAGCTCACTGCAACCTCTGCTCCC 3850
Qy 1350 AGGTTCAAGCAATTTCTCTGCTCTCAGCTCCCGGTAGCTGGGACTACAGCGC----- 1402
Db 3849 AGGTTCAAGTAAATTTCTCTGCTCTCAGCTCCCGGTAGCTGGGATTAAGTGTGTGAT 1521
Qy 1403 -CAGCCCGGCTAAATTTTGTATTTAGTGTAGATGGGGTTTCAACATATTAGCCCGGC 1461
Db 3789 CCACATAGGCTAAATTTTGTATTTAGTGTAGATGGGGTTTCAACATATTAGCCCGGC 3730
Qy 1462 TGGTCTTGAATCTCTGACCTCAGGTGATCCACCACTCAGCTCTCTAAAGTGTGGAT 1521
Db 3729 TGGTCTTGAATCTCTGACCTCAGGTGATCCCGCTCGCTCTCAAAAGTGTGGAT 3670
Qy 1522 TACAGGCATGAGTCAACCGCGCCGCCAAGGTCAGTGTGTTAAAGGAATAACTTCA 1579
Db 3669 TACAGGCATGAGCCTATGTTAGTGTAGATGGGGTTTGGCCATGTTAGCCAGGC 3730

RESULT 14

Qy 1462 TGGTCTTGAATCTCTGACCTCAGGTGATCCACCACTCAGCTCTCTAAAGTGTGGAT 1521
Db 3729 TGGTCTTGAATCTCTGACCTCAGGTGATCCCGCTCGCTCTCAAAAGTGTGGAT 3670
Qy 1522 TACAGGCATGAGTCAACCGCGCCGCCAAGGTCAGTGTGTTAAAGGAATAACTTGA 1579
Db 3669 TACAGGCATGAGCCTATGTTAGTGTAGATGGGGTTTGGCCATGTTAGCCAGGC 3612

RESULT 13

US-10-868-184C-11267/c
; Sequence 11267, Application US/10868184C
; GENERAL INFORMATION:

; APPLICANT: Rosen, et. al
; TITLE OF INVENTION: Human Secreted Proteins
; FILE REFERENCE: PS805
; CURRENT APPLICATION NUMBER: US/10/868,184C
; PRIOR FILING DATE: 2004-06-16
; PRIOR APPLICATION NUMBER: 60/278,650
; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: 09/833,245
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US01/11988
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US00/06043
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06012
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06058
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06044
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06059
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06042
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06014
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13046
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 11267
; LENGTH: 6708
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-868-184C-11267

Query Match 3.4%; Score 179.2; DB 6; Length 6708;
Best Local Similarity 77.9%; Pred. No. 2.9e-34;
Matches 232; Conservative 0; Mismatches 58; Indels 8; Gaps 1;
Qy 1290 TCTGTGCTTACACCTACCTCTGATGCTCTACACCTGAGCTCACTGCAACCTCTGCTCCC 1349
Db 3909 TCTGTCTCTCAGGCTGGAGTACAGTGGGCACAACTCTCAGCTCACTGCAACCTCTGCTCCC 3850
Qy 1350 AGGTTCAAGCAATTTCTCTGCTCTCAGCTCCCGGTAGCTGGGACTACAGCGC----- 1402
Db 3849 AGGTTCAAGTAAATTTCTCTGCTCTCAGCTCCCGGTAGCTGGGATTAAGTGTGTGAT 3790
Qy 1403 -CAGCCCGGCTAAATTTTGTATTTAGTGTAGATGGGGTTTCAACATATTAGCCCGGC 1461
Db 3789 CCACATAGGCTAAATTTTGTATTTAGTGTAGATGGGGTTTGGCCATGTTAGCCAGGC 3730
Qy 1462 TGGTCTTGAATCTCTGACCTCAGGTGATCCACCACTCAGCTCTCTAAAGTGTGGAT 1521
Db 3729 TGGTCTTGAATCTCTGACCTCAGGTGATCCCGCTCGCTCTCAAAAGTGTGGAT 3670
Qy 1522 TACAGGCATGAGTCAACCGCGCCGCCAAGGTCAGTGTGTTAAAGGAATAACTTCA 1579
Db 3669 TACAGGCATGAGCCTATGTTAGTGTAGATGGGGTTTGGCCATGTTAGCCAGGC 3612

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US-10-868-184C-6360/c
; Sequence 6360, Application US/10868184C
; GENERAL INFORMATION:
; APPLICANT: Rosen, et. al
; TITLE OF INVENTION: Human Secreted Proteins
; FILE REFERENCE: PS805
; CURRENT APPLICATION NUMBER: US/10/868,184C
; CURRENT FILING DATE: 2004-06-16
; PRIOR APPLICATION NUMBER: 60/278,650
; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: 09/833,245
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US01/11988
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US00/06043
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06012
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06058
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06044
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06059
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06042
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06014
; PRIOR FILING DATE: 2000-03-09
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13046
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 6360
; LENGTH: 6708
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-868-184C-6360

Query Match          3.4%; Score 179.2; DB 7; Length 6708;
Best Local Similarity 77.9%; Pred. No. 2.9e-34;
Matches 232; Conservative 0; Mismatches 58; Indels 8; Gaps 1;

Qy 1290 TCTGTGCTTACACCTACCTGATGCTCTACACCTGAGCTGCTGCAACCTCTGCTCC 1349
Db 3909 TCTGTCTCTCAGGCTGGAGTACAGTGGCAATCTCAGCTCAGTCACTGCAACCTCTGCTCC 3850
Qy 1350 AGGTTCAAGCAATTCCTGCTCAGCTCCCGCTAGCTGGGACTACAGGCG 1402
Db 3849 AGGTTCAAGTAATTCCTGCTCAGCTCCCGCTAGCTGGGACTACAGTGGCAACCA 3790
Qy 1403 -CAGCCCGGCTAAATTTTGTATTTAGTAGAGATGGGTTTCCACCATATTAGCTGGGAT 1461
Db 3789 CCACACATGGCTAAATTTTGTATTTAGTAGAGATGGGTTTCCACCATATTAGCTGGGAT 3730
Qy 1462 TGGTCTTGAATCTCTGACCTCAGCTGATGCCACCTCAGCTCCCTTAAAGTCTGGGAT 1521
Db 3729 TGGTCTTGAATCTCTGACCTCAGCTGATGCCACCTCAGCTCCCTTAAAGTCTGGGAT 3670
Qy 1522 TACAGGCATGATGATCCCGCCCGCCAGGCTCAGTGTGTTTAAAGGAATAACTTGA 1579
Db 3669 TACAGGCATGATGATCCCGCCCGCCAGGCTCAGTGTGTTTAAAGGAATAACTTGA 3612

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US-10-868-184C-11267/c
; Sequence 11267, Application US/10868184C
; GENERAL INFORMATION:
; APPLICANT: Rosen, et. al
; TITLE OF INVENTION: Human Secreted Proteins
; FILE REFERENCE: PS805
; CURRENT APPLICATION NUMBER: US/10/868,184C
; CURRENT FILING DATE: 2004-06-16
; PRIOR APPLICATION NUMBER: 60/278,650
; PRIOR FILING DATE: 2001-03-27

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; PRIOR APPLICATION NUMBER: 09/833,245
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US01/11988
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: PCT/US00/06043
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06012
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06058
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06044
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06059
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06042
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: PCT/US00/06014
; PRIOR FILING DATE: 2000-03-09
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13046
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 11267
; LENGTH: 6708
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-868-184C-11267

Query Match          3.4%; Score 179.2; DB 7; Length 6708;
Best Local Similarity 77.9%; Pred. No. 2.9e-34;
Matches 232; Conservative 0; Mismatches 58; Indels 8; Gaps 1;

Qy 1290 TCTGTGCTTACACCTACCTGATGCTCTACACCTGAGCTGCTGCAACCTCTGCTCC 1349
Db 3909 TCTGTCTCTCAGGCTGGAGTACAGTGGCAATCTCAGCTCAGTCACTGCAACCTCTGCTCC 3850
Qy 1350 AGGTTCAAGCAATTCCTGCTCAGCTCCCGCTAGCTGGGACTACAGGCG 1402
Db 3849 AGGTTCAAGTAATTCCTGCTCAGCTCCCGCTAGCTGGGACTACAGTGGCAACCA 3790
Qy 1403 -CAGCCCGGCTAAATTTTGTATTTAGTAGAGATGGGTTTCCACCATATTAGCTGGGAT 1461
Db 3789 CCACACATGGCTAAATTTTGTATTTAGTAGAGATGGGTTTCCACCATATTAGCTGGGAT 3730
Qy 1462 TGGTCTTGAATCTCTGACCTCAGCTGATGCCACCTCAGCTCCCTTAAAGTCTGGGAT 1521
Db 3729 TGGTCTTGAATCTCTGACCTCAGCTGATGCCACCTCAGCTCCCTTAAAGTCTGGGAT 3670
Qy 1522 TACAGGCATGATGATCCCGCCCGCCAGGCTCAGTGTGTTTAAAGGAATAACTTGA 1579
Db 3669 TACAGGCATGATGATCCCGCCCGCCAGGCTCAGTGTGTTTAAAGGAATAACTTGA 3612

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